

SEQUENCE LISTING

<110> Agensys, Inc.
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<120> Nucleic Acids and Corresponding Proteins
Entitled 254P1D6B Useful in Treatment and Detection of
Cancer

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Trp Phe Glu Gly Arg Cys Tyr Leu Val Ser Cys Pro His Lys Glu Asn			
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Cys Glu Pro Lys Lys Met Gly Pro Ile Arg Ser Tyr Leu Thr Phe Val			
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Asp Ile Arg Lys Asp Leu Xaa Phe Leu Gly Lys Asp Trp Gly Leu Glu			
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Asp Trp Gly Leu Leu Pro Gly Ser Glu Gly Ala Phe Asn Ser Ser Val			
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Gly Asp Ser Pro Ala Val Pro Ala Glu Thr Gln Gln Asp Pro Glu Leu			
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His Tyr Leu Asn Glu Ser Ala Ser Thr Pro Ala Pro Lys Leu Pro Glu			
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Arg Ser Val Leu Leu Pro Leu Pro Thr Thr Pro Ser Ser Gly Glu Val			
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Leu Glu Lys Glu Lys Ala Ser Gln Leu Gln Glu Gln Ser Ser Asn Ser			
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Ser	Gly	Lys	Glu	Val	Leu	Met	Pro	Ser	His	Ser	Leu	Pro	Pro	Ala	Ser	265	270	275		
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Leu	Glu	Leu	Ser	Ser	Val	Thr	Val	Glu	Lys	Ser	Pro	Val	Leu	Thr	Val					
acc	ccg	ggg	agt	aca	gag	cac	agc	atc	cca	aca	cct	ccc	act	agc	gca	300	305	310		1444
Thr	Pro	Gly	Ser	Thr	Glu	His	Ser	Ile	Pro	Thr	Pro	Pro	Thr	Ser	Ala					
gcc	ccc	tct	gag	tcc	acc	cca	tct	gag	cta	ccc	ata	tct	cct	acc	act	315	320	325		1492
Ala	Pro	Ser	Glu	Ser	Thr	Pro	Ser	Glu	Leu	Pro	Ile	Ser	Pro	Thr	Thr					
gct	ccc	agg	aca	gtg	aaa	gaa	ctt	acg	gta	tcg	gct	gga	gat	aac	cta	330	335	340		1540
Ala	Pro	Arg	Thr	Val	Lys	Glu	Leu	Thr	Val	Ser	Ala	Gly	Asp	Asn	Leu					
att	ata	act	tta	ccc	gac	aat	gaa	gtt	gaa	ctg	aag	gcc	ttt	gtt	gcg	345	350	355		1588
Ile	Ile	Thr	Leu	Pro	Asp	Asn	Glu	Val	Glu	Leu	Lys	Ala	Phe	Val	Ala					
cca	gcg	cca	cct	gta	gaa	aca	acc	tac	aac	tat	gaa	tgg	aat	tta	ata	360	365	370	375	1636
Pro	Ala	Pro	Pro	Val	Glu	Thr	Thr	Tyr	Asn	Tyr	Glu	Trp	Asn	Leu	Ile					
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Ser	His	Pro	Thr	Asp	Tyr	Gln	Gly	Glu	Ile	Lys	Gln	Gly	His	Lys	Gln					
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Thr	Leu	Asn	Leu	Ser	Gln	Leu	Ser	Val	Gly	Leu	Tyr	Val	Phe	Lys	Val					
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Thr	Val	Ser	Ser	Glu	Asn	Ala	Phe	Gly	Glu	Gly	Phe	Val	Asn	Val	Thr					
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Val	Lys	Pro	Ala	Arg	Arg	Val	Asn	Leu	Pro	Pro	Val	Ala	Val	Val	Ser					
ccc	caa	ctg	caa	gag	ctc	act	ttg	cct	ttg	acg	tca	gcc	ctc	att	gtat	440	445	450	455	1876
Pro	Gln	Leu	Gln	Glu	Leu	Thr	Leu	Pro	Leu	Thr	Ser	Ala	Leu	Ile	Asp					
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Glu	Ile	Asn	Gly	Pro	Phe	Ile	Glu	Glu	Lys	Thr	Ser	Val	Asp	Ser	Pro					
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Val	Leu	Arg	Leu	Ser	Asn	Leu	Asp	Pro	Gly	Asn	Tyr	Ser	Phe	Arg	Leu					
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Thr	Val	Thr	Asp	Ser	Asp	Gly	Ala	Thr	Asn	Ser	Thr	Thr	Ala	Ala	Leu					
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Asn His Thr Ile Thr Leu Pro Gln Asn Ser Ile Thr Leu Asn Gly Asn			
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Gln Ser Ser Asp Asp His Gln Ile Val Leu Tyr Glu Trp Ser Leu Gly			
555	560	565	
cct ggg agt gag ggc aaa cat gtg gtc atg cag gga gta cag acg cca			2260
Pro Gly Ser Glu Gly Lys His Val Val Met Gln Gly Val Gln Thr Pro			
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Tyr Leu His Leu Ser Ala Met Gln Glu Gly Asp Tyr Thr Phe Gln Leu			
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aag gtg aca gat tct tca agg caa cag tct act gct gtr gtg act gtg			2356
Lys Val Thr Asp Ser Ser Arg Gln Gln Ser Thr Ala Xaa Val Thr Val			
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att gtc cag cct gaa aac aat aga cct cca gtg gct gtg gcc ggc cct			2404
Ile Val Gln Pro Glu Asn Asn Arg Pro Pro Val Ala Val Ala Gly Pro			
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Asp Lys Glu Leu Ile Phe Pro Val Glu Ser Ala Thr Leu Asp Gly Ser			
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Ser Ser Asp Asp His Gly Ile Val Phe Tyr His Trp Glu His Val			
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Arg Gly Pro Ser Ala Val Glu Met Glu Asn Ile Asp Lys Ala Ile Ala			
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Thr Val Thr Gly Leu Gln Val Gly Thr Tyr His Phe Arg Leu Thr Val			
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Lys Lys Glu Asn Asn Ser Pro Pro Arg Ala Arg Ala Gly Gly Arg His			
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Val Leu Val Leu Pro Asn Asn Ser Ile Thr Leu Asp Gly Ser Arg Ser			
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Thr Asp Asp Gln Arg Ile Val Ser Tyr Leu Trp Ile Arg Asp Gly Gln			
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agt cca gca gct gga gat gtc atc gat ggc tct gac cac agt gtg gct			2836
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Gln	Ser	Arg	Pro	Pro	Phe	Lys	Val	Leu	Lys	Ala	Ala	Glu	Val	Ala	Arg		
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Cys	Glu	Trp	Ser	Ile	Phe	Tyr	Val	Thr	Val	Leu	Ala	Phe	Thr	Leu	Ile		
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gtg	cta	aca	gga	ggt	ttc	act	tgg	ctt	tgc	atc	tgc	tgc	tgc	aaa	aga	3460	
Val	Leu	Thr	Gly	Gly	Phe	Thr	Trp	Leu	Cys	Ile	Cys	Cys	Cys	Lys	Arg		
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caa	aaa	agg	act	aaa	atc	agg	aaa	aaa	aca	aag	tac	acc	atc	ctg	gat	3508	
Gln	Lys	Arg	Thr	Lys	Ile	Arg	Lys	Lys	Thr	Lys	Tyr	Thr	Ile	Leu	Asp		
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Asn	Met	Asp	Glu	Gln	Glu	Arg	Met	Glu	Leu	Arg	Pro	Lys	Tyr	Gly	Ile		
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Leu	Leu	Asp	Tyr	Gly	Asp	Met	Met	Leu	Asn	Arg	Gly	Ser	Pro	Ser	Gly
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Arg	Gly	Ser	Ala	Glu	Tyr	Thr	Asp	Trp	Gly	Leu	Leu	Pro	Gly	Ser	Glu
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Gly	Ala	Phe	Asn	Ser	Ser	Val	Gly	Asp	Ser	Pro	Ala	Val	Pro	Ala	Glu
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Thr	Gln	Gln	Asp	Pro	Glu	Leu	His	Tyr	Leu	Asn	Glu	Ser	Ala	Ser	Thr
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Gln	Glu	Gln	Ser	Ser	Asn	Ser	Ser	Gly	Lys	Glu	Val	Leu	Met	Pro	Ser
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His	Ser	Leu	Pro	Pro	Ala	Ser	Leu	Glu	Leu	Ser	Ser	Val	Thr	Val	Glu
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Lys	Ser	Pro	Val	Leu	Thr	Val	Thr	Pro	Gly	Ser	Thr	Glu	His	Ser	Ile
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Val	Ser	Ala	Gly	Asp	Asn	Leu	Ile	Ile	Thr	Leu	Pro	Asp	Asn	Glu	Val
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Glu	Leu	Lys	Ala	Phe	Val	Ala	Pro	Ala	Pro	Pro	Val	Glu	Thr	Thr	Tyr
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Asn Tyr Glu Trp Asn Leu Ile Ser His Pro Thr Asp Tyr Gln Gly Glu
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 Gly Leu Tyr Val Phe Lys Val Thr Val Ser Ser Glu Asn Ala Phe Gly
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 Glu Gly Phe Val Asn Val Thr Val Lys Pro Ala Arg Arg Val Asn Leu
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 Pro Pro Val Ala Val Val Ser Pro Gln Leu Gln Glu Leu Thr Leu Pro
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 Leu Thr Ser Ala Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr Glu
 450 455 460
 Ile Val Ser Tyr His Trp Glu Glu Ile Asn Gly Pro Phe Ile Glu Glu
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Ala Arg Lys Gln Cys Ser Glu Gly Arg Thr Tyr Ser Asn Ala Val Ile
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Ser Pro Asn Leu Glu Thr Thr Arg Ile Met Arg Val Ser His Thr Phe
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cct gtc gta gac tgc acg gcc gct tgc tgt gac ctg tcc agc tgt gac 915
Pro Val Val Asp Cys Thr Ala Ala Cys Cys Asp Leu Ser Ser Cys Asp
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Leu Ala Trp Trp Phe Glu Gly Arg Cys Tyr Leu Val Ser Cys Pro His
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Lys Glu Asn Cys Glu Pro Lys Met Gly Pro Ile Arg Ser Tyr Leu
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ctc att gat ggc agc caa agt aca gat gat act gaa ata gtg agt tat		2115
Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr Glu Ile Val Ser Tyr		
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cat tgg gaa gaa ata aac ggg ccc ttc ata gaa gag aag act tca gtt		2163
His Trp Glu Glu Ile Asn Gly Pro Phe Ile Glu Glu Lys Thr Ser Val		
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Asp Ser Pro Val Leu Arg Leu Ser Asn Leu Asp Pro Gly Asn Tyr Ser		
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Phe Arg Ile Thr Val Thr Asp Ser Asp Gly Ala Thr Asn Ser Thr Thr		
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Ala Ala Leu Ile Val Asn Asn Ala Val Asp Tyr Pro Pro Val Ala Asn		
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Ser Leu Gly Pro Gly Ser Glu Gly Lys His Val Val Met Gln Gly Val		
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Gln Thr Pro Tyr Leu His Leu Ser Ala Met Gln Glu Gly Asp Tyr Thr		
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Phe Gln Leu Lys Val Thr Asp Ser Ser Arg Gln Gln Ser Thr Ala Val		
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Val Thr Val Ile Val Gln Pro Glu Asn Asn Arg Pro Pro Val Ala Val		
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Ala Gly Pro Asp Lys Glu Leu Ile Phe Pro Val Glu Ser Ala Thr Leu		
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Asp Gly Ser Ser Ser Asp Asp His Gly Ile Val Phe Tyr His Trp		
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Glu His Val Arg Gly Pro Ser Ala Val Glu Met Glu Asn Ile Asp Lys		
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ggc aga cat gtt ctt gtg ctt ccc aat aat tcc att act ttg gat ggt		2931
Gly Arg His Val Leu Val Pro Asn Asn Ser Ile Thr Leu Asp Gly		
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tca agg tct act gat gac caa aga att gtg tcc tat ctg tgg atc cg ^g		2979
Ser Arg Ser Thr Asp Asp Gln Arg Ile Val Ser Tyr Leu Trp Ile Arg		
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Asp Gly Gln Ser Pro Ala Ala Gly Asp Val Ile Asp Gly Ser Asp His		
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Ser Val Ala Leu Gln Leu Thr Asn Leu Val Glu Gly Val Tyr Thr Phe		
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cac ttg cga gtc acc gac agt cag ggg gcc tcg gac aca gac act gcc		3123
His Leu Arg Val Thr Asp Ser Gln Gly Ala Ser Asp Thr Asp Thr Ala		
780	785	790
795		
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Thr Val Glu Val Gln Pro Asp Pro Arg Lys Ser Gly Leu Val Glu Leu		
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Thr Leu Gln Val Gly Val Gly Gln Leu Thr Glu Gln Arg Lys Asp Thr		
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Leu Val Arg Gln Leu Ala Val Leu Leu Asn Val Leu Asp Ser Asp Ile		
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Lys Val Gln Lys Ile Arg Ala His Ser Asp Leu Ser Thr Val Ile Val		
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Phe Tyr Val Gln Ser Arg Pro Pro Phe Lys Val Leu Lys Ala Ala Glu		
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gtg gcc cga aat ctg cac atg cg ^g ctc tca aag gag aag gct gac ttc		3411
Val Ala Arg Asn Leu His Met Arg Leu Ser Lys Glu Lys Ala Asp Phe		
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Leu Leu Phe Lys Val Leu Arg Val Asp Thr Ala Gly Cys Leu Leu Lys		
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Cys Ser Gly His Gly His Cys Asp Pro Leu Thr Lys Arg Cys Ile Cys		
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Thr	Leu	Ile	Val	Leu	Thr	Gly	Gly	Phe	Thr	Trp	Leu	Cys	Ile	Cys	Cys	
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 50 55 60
 Glu Gly Arg Cys Tyr Leu Val Ser Cys Pro His Lys Glu Asn Cys Glu
 65 70 75 80
 Pro Lys Lys Met Gly Pro Ile Arg Ser Tyr Leu Thr Phe Val Leu Arg
 85 90 95
 Pro Val Gln Arg Pro Ala Gln Leu Leu Asp Tyr Gly Asp Met Met Leu
 100 105 110
 Asn Arg Gly Ser Pro Ser Gly Ile Trp Gly Asp Ser Pro Glu Asp Ile
 115 120 125
 Arg Lys Asp Leu Pro Phe Leu Gly Lys Asp Trp Gly Leu Glu Glu Met
 130 135 140
 Ser Glu Tyr Ser Asp Asp Tyr Arg Glu Leu Glu Lys Asp Leu Leu Gln
 145 150 155 160
 Pro Ser Gly Lys Gln Glu Pro Arg Gly Ser Ala Glu Tyr Thr Asp Trp
 165 170 175
 Gly Leu Leu Pro Gly Ser Glu Gly Ala Phe Asn Ser Ser Val Gly Asp
 180 185 190
 Ser Pro Ala Val Pro Ala Glu Thr Gln Gln Asp Pro Glu Leu His Tyr
 195 200 205
 Leu Asn Glu Ser Ala Ser Thr Pro Ala Pro Lys Leu Pro Glu Arg Ser
 210 215 220
 Val Leu Leu Pro Leu Pro Thr Thr Pro Ser Ser Gly Glu Val Leu Glu
 225 230 235 240
 Lys Glu Lys Ala Ser Gln Leu Gln Glu Gln Ser Ser Asn Ser Ser Gly
 245 250 255
 Lys Glu Val Leu Met Pro Ser His Ser Leu Pro Pro Ala Ser Leu Glu
 260 265 270
 Leu Ser Ser Val Thr Val Glu Lys Ser Pro Val Leu Thr Val Thr Pro
 275 280 285
 Gly Ser Thr Glu His Ser Ile Pro Thr Pro Pro Thr Ser Ala Ala Pro
 290 295 300

Ser Glu Ser Thr Pro Ser Glu Leu Pro Ile Ser Pro Thr Thr Ala Pro
 305 310 315 320
 Arg Thr Val Lys Glu Leu Thr Val Ser Ala Gly Asp Asn Leu Ile Ile
 325 330 335
 Thr Leu Pro Asp Asn Glu Val Glu Leu Lys Ala Phe Val Ala Pro Ala
 340 345 350
 Pro Pro Val Glu Thr Thr Tyr Asn Tyr Glu Trp Asn Leu Ile Ser His
 355 360 365
 Pro Thr Asp Tyr Gln Gly Glu Ile Lys Gln Gly His Lys Gln Thr Leu
 370 375 380
 Asn Leu Ser Gln Leu Ser Val Gly Leu Tyr Val Phe Lys Val Thr Val
 385 390 395 400
 Ser Ser Glu Asn Ala Phe Gly Glu Gly Phe Val Asn Val Thr Val Lys
 405 410 415
 Pro Ala Arg Arg Val Asn Leu Pro Pro Val Ala Val Val Ser Pro Gln
 420 425 430
 Leu Gln Glu Leu Thr Leu Pro Leu Thr Ser Ala Leu Ile Asp Gly Ser
 435 440 445
 Gln Ser Thr Asp Asp Thr Glu Ile Val Ser Tyr His Trp Glu Glu Ile
 450 455 460
 Asn Gly Pro Phe Ile Glu Glu Lys Thr Ser Val Asp Ser Pro Val Leu
 465 470 475 480
 Arg Leu Ser Asn Leu Asp Pro Gly Asn Tyr Ser Phe Arg Leu Thr Val
 485 490 495
 Thr Asp Ser Asp Gly Ala Thr Asn Ser Thr Thr Ala Ala Leu Ile Val
 500 505 510
 Asn Asn Ala Val Asp Tyr Pro Pro Val Ala Asn Ala Gly Pro Asn His
 515 520 525
 Thr Ile Thr Leu Pro Gln Asn Ser Ile Thr Leu Asn Gly Asn Gln Ser
 530 535 540
 Ser Asp Asp His Gln Ile Val Leu Tyr Glu Trp Ser Leu Gly Pro Gly
 545 550 555 560
 Ser Glu Gly Lys His Val Val Met Gln Gly Val Gln Thr Pro Tyr Leu
 565 570 575
 His Leu Ser Ala Met Gln Glu Gly Asp Tyr Thr Phe Gln Leu Lys Val
 580 585 590
 Thr Asp Ser Ser Arg Gln Gln Ser Thr Ala Val Val Thr Val Ile Val
 595 600 605
 Gln Pro Glu Asn Asn Arg Pro Pro Val Ala Val Ala Gly Pro Asp Lys
 610 615 620
 Glu Leu Ile Phe Pro Val Glu Ser Ala Thr Leu Asp Gly Ser Ser Ser
 625 630 635 640
 Ser Asp Asp His Gly Ile Val Phe Tyr His Trp Glu His Val Arg Gly
 645 650 655
 Pro Ser Ala Val Glu Met Glu Asn Ile Asp Lys Ala Ile Ala Thr Val
 660 665 670
 Thr Gly Leu Gln Val Gly Thr Tyr His Phe Arg Leu Thr Val Lys Asp
 675 680 685
 Gln Gln Gly Leu Ser Ser Thr Ser Thr Leu Thr Val Ala Val Lys Lys
 690 695 700
 Glu Asn Asn Ser Pro Pro Arg Ala Arg Ala Gly Gly Arg His Val Leu
 705 710 715 720
 Val Leu Pro Asn Asn Ser Ile Thr Leu Asp Gly Ser Arg Ser Thr Asp
 725 730 735
 Asp Gln Arg Ile Val Ser Tyr Leu Trp Ile Arg Asp Gly Gln Ser Pro
 740 745 750
 Ala Ala Gly Asp Val Ile Asp Gly Ser Asp His Ser Val Ala Leu Gln
 755 760 765
 Leu Thr Asn Leu Val Glu Gly Val Tyr Thr Phe His Leu Arg Val Thr
 770 775 780
 Asp Ser Gln Gly Ala Ser Asp Thr Asp Thr Ala Thr Val Glu Val Gln
 785 790 795 800
 Pro Asp Pro Arg Lys Ser Gly Leu Val Glu Leu Thr Leu Gln Val Gly
 805 810 815

Val Gly Gln Leu Thr Glu Gln Arg Lys Asp Thr Leu Val Arg Gln Leu
 820 825 830
 Ala Val Leu Leu Asn Val Leu Asp Ser Asp Ile Lys Val Gln Lys Ile
 835 840 845
 Arg Ala His Ser Asp Leu Ser Thr Val Ile Val Phe Tyr Val Gln Ser
 850 855 860
 Arg Pro Pro Phe Lys Val Leu Lys Ala Ala Glu Val Ala Arg Asn Leu
 865 870 875 880
 His Met Arg Leu Ser Lys Glu Lys Ala Asp Phe Leu Leu Phe Lys Val
 885 890 895
 Leu Arg Val Asp Thr Ala Gly Cys Leu Leu Lys Cys Ser Gly His Gly
 900 905 910
 His Cys Asp Pro Leu Thr Lys Arg Cys Ile Cys Ser His Leu Trp Met
 915 920 925
 Glu Asn Leu Ile Gln Arg Tyr Ile Trp Asp Gly Glu Ser Asn Cys Glu
 930 935 940
 Trp Ser Ile Phe Tyr Val Thr Val Leu Ala Phe Thr Leu Ile Val Leu
 945 950 955 960
 Thr Gly Gly Phe Thr Trp Leu Cys Ile Cys Cys Cys Lys Arg Gln Lys
 965 970 975
 Arg Thr Lys Ile Arg Lys Lys Thr Lys Tyr Thr Ile Leu Asp Asn Met
 980 985 990
 Asp Glu Gln Glu Arg Met Glu Leu Arg Pro Lys Tyr Gly Ile Lys His
 995 1000 1005
 Arg Ser Thr Glu His Asn Ser Ser Leu Met Val Ser Glu Ser Glu Phe
 1010 1015 1020
 Asp Ser Asp Gln Asp Thr Ile Phe Ser Arg Glu Lys Met Glu Arg Gly
 1025 1030 1035 1040
 Asn Pro Lys Val Ser Met Asn Gly Ser Ile Arg Asn Gly Ala Ser Phe
 1045 1050 1055
 Ser Tyr Cys Ser Lys Asp Arg
 1060

<210> 8
 <211> 1072
 <212> PRT
 <213> Homo sapiens

<400> 8
 Met Ala Pro Pro Thr Gly Val Leu Ser Ser Leu Leu Leu Val Thr
 1 5 10 15
 Ile Ala Gly Cys Ala Arg Lys Gln Cys Ser Glu Gly Arg Thr Tyr Ser
 20 25 30
 Asn Ala Val Ile Ser Pro Asn Leu Glu Thr Thr Arg Ile Met Arg Val
 35 40 45
 Ser His Thr Phe Pro Val Val Asp Cys Thr Ala Ala Cys Cys Asp Leu
 50 55 60
 Ser Ser Cys Asp Leu Ala Trp Trp Phe Glu Gly Arg Cys Tyr Leu Val
 65 70 75 80
 Ser Cys Pro His Lys Glu Asn Cys Glu Pro Lys Lys Met Gly Pro Ile
 85 90 95
 Arg Ser Tyr Leu Thr Phe Val Leu Arg Pro Val Gln Arg Pro Ala Gln
 100 105 110
 Leu Leu Asp Tyr Gly Asp Met Met Leu Asn Arg Gly Ser Pro Ser Gly
 115 120 125
 Ile Trp Gly Asp Ser Pro Glu Asp Ile Arg Lys Asp Leu Pro Phe Leu
 130 135 140
 Gly Lys Asp Trp Gly Leu Glu Glu Met Ser Glu Tyr Ser Asp Asp Tyr
 145 150 155 160
 Arg Glu Leu Glu Lys Asp Leu Leu Gln Pro Ser Gly Lys Gln Glu Pro
 165 170 175
 Arg Gly Ser Ala Glu Tyr Thr Asp Trp Gly Leu Leu Pro Gly Ser Glu
 180 185 190

Gly Ala Phe Asn Ser Ser Val Gly Asp Ser Pro Ala Val Pro Ala Glu
 195 200 205
 Thr Gln Gln Asp Pro Glu Leu His Tyr Leu Asn Glu Ser Ala Ser Thr
 210 215 220
 Pro Ala Pro Lys Leu Pro Glu Arg Ser Val Leu Leu Pro Leu Pro Thr
 225 230 235 240
 Thr Pro Ser Ser Gly Glu Val Leu Glu Lys Glu Lys Ala Ser Gln Leu
 245 250 255
 Gln Glu Gln Ser Ser Asn Ser Ser Gly Lys Glu Val Leu Met Pro Ser
 260 265 270
 His Ser Leu Pro Pro Ala Ser Leu Glu Leu Ser Ser Val Thr Val Glu
 275 280 285
 Lys Ser Pro Val Leu Thr Val Thr Pro Gly Ser Thr Glu His Ser Ile
 290 295 300
 Pro Thr Pro Pro Thr Ser Ala Ala Pro Ser Glu Ser Thr Pro Ser Glu
 305 310 315 320
 Leu Pro Ile Ser Pro Thr Thr Ala Pro Arg Thr Val Lys Glu Leu Thr
 325 330 335
 Val Ser Ala Gly Asp Asn Leu Ile Ile Thr Leu Pro Asp Asn Glu Val
 340 345 350
 Glu Leu Lys Ala Phe Val Ala Pro Ala Pro Pro Val Glu Thr Thr Tyr
 355 360 365
 Asn Tyr Glu Trp Asn Leu Ile Ser His Pro Thr Asp Tyr Gln Gly Glu
 370 375 380
 Ile Lys Gln Gly His Lys Gln Thr Leu Asn Leu Ser Gln Leu Ser Val
 385 390 395 400
 Gly Leu Tyr Val Phe Lys Val Thr Val Ser Ser Glu Asn Ala Phe Gly
 405 410 415
 Glu Gly Phe Val Asn Val Thr Val Lys Pro Ala Arg Arg Val Asn Leu
 420 425 430
 Pro Pro Val Ala Val Val Ser Pro Gln Leu Gln Glu Leu Thr Leu Pro
 435 440 445
 Leu Thr Ser Ala Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr Glu
 450 455 460
 Ile Val Ser Tyr His Trp Glu Glu Ile Asn Gly Pro Phe Ile Glu Glu
 465 470 475 480
 Lys Thr Ser Val Asp Ser Pro Val Leu Arg Leu Ser Asn Leu Asp Pro
 485 490 495
 Gly Asn Tyr Ser Phe Arg Leu Thr Val Thr Asp Ser Asp Gly Ala Thr
 500 505 510
 Asn Ser Thr Thr Ala Ala Leu Ile Val Asn Asn Ala Val Asp Tyr Pro
 515 520 525
 Pro Val Ala Asn Ala Gly Pro Asn His Thr Ile Thr Leu Pro Gln Asn
 530 535 540
 Ser Ile Thr Leu Asn Gly Asn Gln Ser Ser Asp Asp His Gln Ile Val
 545 550 555 560
 Leu Tyr Glu Trp Ser Leu Gly Pro Gly Ser Glu Gly Lys His Val Val
 565 570 575
 Met Gln Gly Val Gln Thr Pro Tyr Leu His Leu Ser Ala Met Gln Glu
 580 585 590
 Gly Asp Tyr Thr Phe Gln Leu Lys Val Thr Asp Ser Ser Arg Gln Gln
 595 600 605
 Ser Thr Ala Val Val Thr Val Ile Val Gln Pro Glu Asn Asn Arg Pro
 610 615 620
 Pro Val Ala Val Ala Gly Pro Asp Lys Glu Leu Ile Phe Pro Val Glu
 625 630 635 640
 Ser Ala Thr Leu Asp Gly Ser Ser Ser Asp Asp His Gly Ile Val
 645 650 655
 Phe Tyr His Trp Glu His Val Arg Gly Pro Ser Ala Val Glu Met Glu
 660 665 670
 Asn Ile Asp Lys Ala Ile Ala Thr Val Thr Gly Leu Gln Val Gly Thr
 675 680 685
 Tyr His Phe Arg Leu Thr Val Lys Asp Gln Gln Gly Leu Ser Ser Thr
 690 695 700

Ser Thr Leu Thr Val Ala Val Lys Lys Glu Asn Asn Ser Pro Pro Arg
 705 710 715 720
 Ala Arg Ala Gly Gly Arg His Val Leu Val Leu Pro Asn Asn Ser Ile
 725 730 735
 Thr Leu Asp Gly Ser Arg Ser Thr Asp Asp Gln Arg Ile Val Ser Tyr
 740 745 750
 Leu Trp Ile Arg Asp Gly Gln Ser Pro Ala Ala Gly Asp Val Ile Asp
 755 760 765
 Gly Ser Asp His Ser Val Ala Leu Gln Leu Thr Asn Leu Val Glu Gly
 770 775 780
 Val Tyr Thr Phe His Leu Arg Val Thr Asp Ser Gln Gly Ala Ser Asp
 785 790 795 800
 Thr Asp Thr Ala Thr Val Glu Val Gln Pro Asp Pro Arg Lys Ser Gly
 805 810 815
 Leu Val Glu Leu Thr Leu Gln Val Gly Val Gly Gln Leu Thr Glu Gln
 820 825 830
 Arg Lys Asp Thr Leu Val Arg Gln Leu Ala Val Leu Leu Asn Val Leu
 835 840 845
 Asp Ser Asp Ile Lys Val Gln Lys Ile Arg Ala His Ser Asp Leu Ser
 850 855 860
 Thr Val Ile Val Phe Tyr Val Gln Ser Arg Pro Pro Phe Lys Val Leu
 865 870 875 880
 Lys Ala Ala Glu Val Ala Arg Asn Leu His Met Arg Leu Ser Lys Glu
 885 890 895
 Lys Ala Asp Phe Leu Leu Phe Lys Val Leu Arg Val Asp Thr Ala Gly
 900 905 910
 Cys Leu Leu Lys Cys Ser Gly His Gly His Cys Asp Pro Leu Thr Lys
 915 920 925
 Arg Cys Ile Cys Ser His Leu Trp Met Glu Asn Leu Ile Gln Arg Tyr
 930 935 940
 Ile Trp Asp Gly Glu Ser Asn Cys Glu Trp Ser Ile Phe Tyr Val Thr
 945 950 955 960
 Val Leu Ala Phe Thr Leu Ile Val Leu Thr Gly Gly Phe Thr Trp Leu
 965 970 975
 Cys Ile Cys Cys Lys Arg Gln Lys Arg Thr Lys Ile Arg Lys Lys
 980 985 990
 Thr Lys Tyr Thr Ile Leu Asp Asn Met Asp Glu Gln Glu Arg Met Glu
 995 1000 1005
 Leu Arg Pro Lys Tyr Gly Ile Lys His Arg Ser Thr Glu His Asn Ser
 1010 1015 1020
 Ser Leu Met Val Ser Glu Ser Glu Phe Asp Ser Asp Gln Asp Thr Ile
 1025 1030 1035 1040
 Phe Ser Arg Glu Lys Met Glu Arg Gly Asn Pro Lys Val Ser Met Asn
 1045 1050 1055
 Gly Ser Ile Arg Asn Gly Ala Ser Phe Ser Tyr Cys Ser Lys Asp Arg
 1060 1065 1070

<210> 9
 <211> 1072
 <212> PRT
 <213> Homo sapiens

<400> 9
 Met Ala Pro Pro Thr Gly Val Leu Ser Ser Leu Leu Leu Val Thr
 1 5 10 15
 Ile Ala Gly Cys Ala Arg Lys Gln Cys Ser Glu Gly Arg Thr Tyr Ser
 20 25 30
 Asn Ala Val Ile Ser Pro Asn Leu Glu Thr Thr Arg Ile Met Arg Val
 35 40 45
 Ser His Thr Phe Pro Val Val Asp Cys Thr Ala Ala Cys Cys Asp Leu
 50 55 60
 Ser Ser Cys Asp Leu Ala Trp Trp Phe Glu Gly Arg Cys Tyr Leu Val
 65 70 75 80

Ser Cys Pro His Lys Glu Asn Cys Glu Pro Lys Lys Met Gly Pro Ile
 85 90 95
 Arg Ser Tyr Leu Thr Phe Val Leu Arg Pro Val Gln Arg Pro Ala Gln
 100 105 110
 Leu Leu Asp Tyr Gly Asp Met Met Leu Asn Arg Gly Ser Pro Ser Gly
 115 120 125
 Ile Trp Gly Asp Ser Pro Glu Asp Ile Arg Lys Asp Leu Pro Phe Leu
 130 135 140
 Gly Lys Asp Trp Gly Leu Glu Glu Met Ser Glu Tyr Ala Asp Asp Tyr
 145 150 155 160
 Arg Glu Leu Glu Lys Asp Leu Leu Gln Pro Ser Gly Lys Gln Glu Pro
 165 170 175
 Arg Gly Ser Ala Glu Tyr Thr Asp Trp Gly Leu Leu Pro Gly Ser Glu
 180 185 190
 Gly Ala Phe Asn Ser Ser Val Gly Asp Ser Pro Ala Val Pro Ala Glu
 195 200 205
 Thr Gln Gln Asp Pro Glu Leu His Tyr Leu Asn Glu Ser Ala Ser Thr
 210 215 220
 Pro Ala Pro Lys Leu Pro Glu Arg Ser Val Leu Leu Pro Leu Pro Thr
 225 230 235 240
 Thr Pro Ser Ser Gly Glu Val Leu Glu Lys Glu Lys Ala Ser Gln Leu
 245 250 255
 Gln Glu Gln Ser Ser Asn Ser Ser Gly Lys Glu Val Leu Met Pro Ser
 260 265 270
 His Ser Leu Pro Pro Ala Ser Leu Glu Leu Ser Ser Val Thr Val Glu
 275 280 285
 Lys Ser Pro Val Leu Thr Val Thr Pro Gly Ser Thr Glu His Ser Ile
 290 295 300
 Pro Thr Pro Pro Thr Ser Ala Ala Pro Ser Glu Ser Thr Pro Ser Glu
 305 310 315 320
 Leu Pro Ile Ser Pro Thr Thr Ala Pro Arg Thr Val Lys Glu Leu Thr
 325 330 335
 Val Ser Ala Gly Asp Asn Leu Ile Ile Thr Leu Pro Asp Asn Glu Val
 340 345 350
 Glu Leu Lys Ala Phe Val Ala Pro Ala Pro Pro Val Glu Thr Thr Tyr
 355 360 365
 Asn Tyr Glu Trp Asn Leu Ile Ser His Pro Thr Asp Tyr Gln Gly Glu
 370 375 380
 Ile Lys Gln Gly His Lys Gln Thr Leu Asn Leu Ser Gln Leu Ser Val
 385 390 395 400
 Gly Leu Tyr Val Phe Lys Val Thr Val Ser Ser Glu Asn Ala Phe Gly
 405 410 415
 Glu Gly Phe Val Asn Val Thr Val Lys Pro Ala Arg Arg Val Asn Leu
 420 425 430
 Pro Pro Val Ala Val Val Ser Pro Gln Leu Gln Glu Leu Thr Leu Pro
 435 440 445
 Leu Thr Ser Ala Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr Glu
 450 455 460
 Ile Val Ser Tyr His Trp Glu Glu Ile Asn Gly Pro Phe Ile Glu Glu
 465 470 475 480
 Lys Thr Ser Val Asp Ser Pro Val Leu Arg Leu Ser Asn Leu Asp Pro
 485 490 495
 Gly Asn Tyr Ser Phe Arg Leu Thr Val Thr Asp Ser Asp Gly Ala Thr
 500 505 510
 Asn Ser Thr Thr Ala Ala Leu Ile Val Asn Asn Ala Val Asp Tyr Pro
 515 520 525
 Pro Val Ala Asn Ala Gly Pro Asn His Thr Ile Thr Leu Pro Gln Asn
 530 535 540
 Ser Ile Thr Leu Asn Gly Asn Gln Ser Ser Asp Asp His Gln Ile Val
 545 550 555 560
 Leu Tyr Glu Trp Ser Leu Gly Pro Gly Ser Glu Gly Lys His Val Val
 565 570 575
 Met Gln Gly Val Gln Thr Pro Tyr Leu His Leu Ser Ala Met Gln Glu
 580 585 590

Gly Asp Tyr Thr Phe Gln Leu Lys Val Thr Asp Ser Ser Arg Gln Gln
 595 600 605
 Ser Thr Ala Val Val Thr Val Ile Val Gln Pro Glu Asn Asn Arg Pro
 610 615 620
 Pro Val Ala Val Ala Gly Pro Asp Lys Glu Leu Ile Phe Pro Val Glu
 625 630 635 640
 Ser Ala Thr Leu Asp Gly Ser Ser Ser Asp Asp His Gly Ile Val
 645 650 655
 Phe Tyr His Trp Glu His Val Arg Gly Pro Ser Ala Val Glu Met Glu
 660 665 670
 Asn Ile Asp Lys Ala Ile Ala Thr Val Thr Gly Leu Gln Val Gly Thr
 675 680 685
 Tyr His Phe Arg Leu Thr Val Lys Asp Gln Gln Gly Leu Ser Ser Thr
 690 695 700
 Ser Thr Leu Thr Val Ala Val Lys Lys Glu Asn Asn Ser Pro Pro Arg
 705 710 715 720
 Ala Arg Ala Gly Gly Arg His Val Leu Val Leu Pro Asn Asn Ser Ile
 725 730 735
 Thr Leu Asp Gly Ser Arg Ser Thr Asp Asp Gln Arg Ile Val Ser Tyr
 740 745 750
 Leu Trp Ile Arg Asp Gly Gln Ser Pro Ala Ala Gly Asp Val Ile Asp
 755 760 765
 Gly Ser Asp His Ser Val Ala Leu Gln Leu Thr Asn Leu Val Glu Gly
 770 775 780
 Val Tyr Thr Phe His Leu Arg Val Thr Asp Ser Gln Gly Ala Ser Asp
 785 790 795 800
 Thr Asp Thr Ala Thr Val Glu Val Gln Pro Asp Pro Arg Lys Ser Gly
 805 810 815
 Leu Val Glu Leu Thr Leu Gln Val Gly Val Gly Gln Leu Thr Glu Gln
 820 825 830
 Arg Lys Asp Thr Leu Val Arg Gln Leu Ala Val Leu Leu Asn Val Leu
 835 840 845
 Asp Ser Asp Ile Lys Val Gln Lys Ile Arg Ala His Ser Asp Leu Ser
 850 855 860
 Thr Val Ile Val Phe Tyr Val Gln Ser Arg Pro Pro Phe Lys Val Leu
 865 870 875 880
 Lys Ala Ala Glu Val Ala Arg Asn Leu His Met Arg Leu Ser Lys Glu
 885 890 895
 Lys Ala Asp Phe Leu Leu Phe Lys Val Leu Arg Val Asp Thr Ala Gly
 900 905 910
 Cys Leu Leu Lys Cys Ser Gly His Gly His Cys Asp Pro Leu Thr Lys
 915 920 925
 Arg Cys Ile Cys Ser His Leu Trp Met Glu Asn Leu Ile Gln Arg Tyr
 930 935 940
 Ile Trp Asp Gly Glu Ser Asn Cys Glu Trp Ser Ile Phe Tyr Val Thr
 945 950 955 960
 Val Leu Ala Phe Thr Leu Ile Val Leu Thr Gly Gly Phe Thr Trp Leu
 965 970 975
 Cys Ile Cys Cys Lys Arg Gln Lys Arg Thr Lys Ile Arg Lys Lys
 980 985 990
 Thr Lys Tyr Thr Ile Leu Asp Asn Met Asp Glu Gln Glu Arg Met Glu
 995 1000 1005
 Leu Arg Pro Lys Tyr Gly Ile Lys His Arg Ser Thr Glu His Asn Ser
 1010 1015 1020
 Ser Leu Met Val Ser Glu Ser Glu Phe Asp Ser Asp Gln Asp Thr Ile
 1025 1030 1035 1040
 Phe Ser Arg Glu Lys Met Glu Arg Gly Asn Pro Lys Val Ser Met Asn
 1045 1050 1055
 Gly Ser Ile Arg Asn Gly Ala Ser Phe Ser Tyr Cys Ser Lys Asp Arg
 1060 1065 1070

<210> 10
 <211> 1063

<212> PRT
<213> Homo sapiens

<400> 10
Met Thr Arg Leu Gly Trp Pro Ser Pro Cys Cys Ala Arg Lys Gln Cys
1 5 10 15
Ser Glu Gly Arg Thr Tyr Ser Asn Ala Val Ile Ser Pro Asn Leu Glu
20 25 30
Thr Thr Arg Ile Met Arg Val Ser His Thr Phe Pro Val Val Asp Cys
35 40 45
Thr Ala Ala Cys Cys Asp Leu Ser Ser Cys Asp Leu Ala Trp Trp Phe
50 55 60
Glu Gly Arg Cys Tyr Leu Val Ser Cys Pro His Lys Glu Asn Cys Glu
65 70 75 80
Pro Lys Lys Met Gly Pro Ile Arg Ser Tyr Leu Thr Phe Val Leu Arg
85 90 95
Pro Val Gln Arg Pro Ala Gln Leu Leu Asp Tyr Gly Asp Met Met Leu
100 105 110
Asn Arg Gly Ser Pro Ser Gly Ile Trp Gly Asp Ser Pro Glu Asp Ile
115 120 125
Arg Lys Asp Leu Pro Phe Leu Gly Lys Asp Trp Gly Leu Glu Glu Met
130 135 140
Ser Glu Tyr Ser Asp Asp Tyr Arg Glu Leu Glu Lys Asp Leu Leu Gln
145 150 155 160
Pro Ser Gly Lys Gln Glu Pro Arg Gly Ser Ala Glu Tyr Thr Asp Trp
165 170 175
Gly Leu Leu Pro Gly Ser Glu Gly Ala Phe Asn Ser Ser Val Gly Asp
180 185 190
Ser Pro Ala Val Pro Ala Glu Thr Gln Gln Asp Pro Glu Leu His Tyr
195 200 205
Leu Asn Glu Ser Ala Ser Thr Pro Ala Pro Lys Leu Pro Glu Arg Ser
210 215 220
Val Leu Leu Pro Leu Pro Thr Thr Pro Ser Ser Gly Glu Val Leu Glu
225 230 235 240
Lys Glu Lys Ala Ser Gln Leu Gln Glu Gln Ser Ser Asn Ser Ser Gly
245 250 255
Lys Glu Val Leu Met Pro Ser His Ser Leu Pro Pro Ala Ser Leu Glu
260 265 270
Leu Ser Ser Val Thr Val Glu Lys Ser Pro Val Leu Thr Val Thr Pro
275 280 285
Gly Ser Thr Glu His Ser Ile Pro Thr Pro Pro Thr Ser Ala Ala Pro
290 295 300
Ser Glu Ser Thr Pro Ser Glu Leu Pro Ile Ser Pro Thr Thr Ala Pro
305 310 315 320
Arg Thr Val Lys Glu Leu Thr Val Ser Ala Gly Asp Asn Leu Ile Ile
325 330 335
Thr Leu Pro Asp Asn Glu Val Glu Leu Lys Ala Phe Val Ala Pro Ala
340 345 350
Pro Pro Val Glu Thr Thr Tyr Asn Tyr Glu Trp Asn Leu Ile Ser His
355 360 365
Pro Thr Asp Tyr Gln Gly Glu Ile Lys Gln Gly His Lys Gln Thr Leu
370 375 380
Asn Leu Ser Gln Leu Ser Val Gly Leu Tyr Val Phe Lys Val Thr Val
385 390 395 400
Ser Ser Glu Asn Ala Phe Gly Glu Gly Phe Val Asn Val Thr Val Lys
405 410 415
Pro Ala Arg Arg Val Asn Leu Pro Pro Val Ala Val Val Ser Pro Gln
420 425 430
Leu Gln Glu Leu Thr Leu Pro Leu Thr Ser Ala Leu Ile Asp Gly Ser
435 440 445
Gln Ser Thr Asp Asp Thr Glu Ile Val Ser Tyr His Trp Glu Glu Ile
450 455 460
Asn Gly Pro Phe Ile Glu Glu Lys Thr Ser Val Asp Ser Pro Val Leu
465 470 475 480

Arg Leu Ser Asn Leu Asp Pro Gly Asn Tyr Ser Phe Arg Leu Thr Val
 485 490 495
 Thr Asp Ser Asp Gly Ala Thr Asn Ser Thr Thr Ala Ala Leu Ile Val
 500 505 510
 Asn Asn Ala Val Asp Tyr Pro Pro Val Ala Asn Ala Gly Pro Asn His
 515 520 525
 Thr Ile Thr Leu Pro Gln Asn Ser Ile Thr Leu Asn Gly Asn Gln Ser
 530 535 540
 Ser Asp Asp His Gln Ile Val Leu Tyr Glu Trp Ser Leu Gly Pro Gly
 545 550 555 560
 Ser Glu Gly Lys His Val Val Met Gln Gly Val Gln Thr Pro Tyr Leu
 565 570 575
 His Leu Ser Ala Met Gln Glu Gly Asp Tyr Thr Phe Gln Leu Lys Val
 580 585 590
 Thr Asp Ser Ser Arg Gln Gln Ser Thr Ala Val Val Thr Val Ile Val
 595 600 605
 Gln Pro Glu Asn Asn Arg Pro Pro Val Ala Val Ala Gly Pro Asp Lys
 610 615 620
 Glu Leu Ile Phe Pro Val Glu Ser Ala Thr Leu Asp Gly Ser Ser Ser
 625 630 635 640
 Ser Asp Asp His Gly Ile Val Phe Tyr His Trp Glu His Val Arg Gly
 645 650 655
 Pro Ser Ala Val Glu Met Glu Asn Ile Asp Lys Ala Ile Ala Thr Val
 660 665 670
 Thr Gly Leu Gln Val Gly Thr Tyr His Phe Arg Leu Thr Val Lys Asp
 675 680 685
 Gln Gln Gly Leu Ser Ser Thr Ser Thr Leu Thr Val Ala Val Lys Lys
 690 695 700
 Glu Asn Asn Ser Pro Pro Arg Ala Arg Ala Gly Gly Arg His Val Leu
 705 710 715 720
 Val Leu Pro Asn Asn Ser Ile Thr Leu Asp Gly Ser Arg Ser Thr Asp
 725 730 735
 Asp Gln Arg Ile Val Ser Tyr Leu Trp Ile Arg Asp Gly Gln Ser Pro
 740 745 750
 Ala Ala Gly Asp Val Ile Asp Gly Ser Asp His Ser Val Ala Leu Gln
 755 760 765
 Leu Thr Asn Leu Val Glu Gly Val Tyr Thr Phe His Leu Arg Val Thr
 770 775 780
 Asp Ser Gln Gly Ala Ser Asp Thr Asp Thr Ala Thr Val Glu Val Gln
 785 790 795 800
 Pro Asp Pro Arg Lys Ser Gly Leu Val Glu Leu Thr Leu Gln Val Gly
 805 810 815
 Val Gly Gln Leu Thr Glu Gln Arg Lys Asp Thr Leu Val Arg Gln Leu
 820 825 830
 Ala Val Leu Leu Asn Val Leu Asp Ser Asp Ile Lys Val Gln Lys Ile
 835 840 845
 Arg Ala His Ser Asp Leu Ser Thr Val Ile Val Phe Tyr Val Gln Ser
 850 855 860
 Arg Pro Pro Phe Lys Val Leu Lys Ala Ala Glu Val Ala Arg Asn Leu
 865 870 875 880
 His Met Arg Leu Ser Lys Glu Lys Ala Asp Phe Leu Leu Phe Lys Val
 885 890 895
 Leu Arg Val Asp Thr Ala Gly Cys Leu Leu Lys Cys Ser Gly His Gly
 900 905 910
 His Cys Asp Pro Leu Thr Lys Arg Cys Ile Cys Ser His Leu Trp Met
 915 920 925
 Glu Asn Leu Ile Gln Arg Tyr Ile Trp Asp Gly Glu Ser Asn Cys Glu
 930 935 940
 Trp Ser Ile Phe Tyr Val Thr Val Leu Ala Phe Thr Leu Ile Val Leu
 945 950 955 960
 Thr Gly Gly Phe Thr Trp Leu Cys Ile Cys Cys Cys Lys Arg Gln Lys
 965 970 975
 Arg Thr Lys Ile Arg Lys Lys Thr Lys Tyr Thr Ile Leu Asp Asn Met
 980 985 990

Asp Glu Gln Glu Arg Met Glu Leu Arg Pro Lys Tyr Gly Ile Lys His
 995 1000 1005
 Arg Ser Thr Glu His Asn Ser Ser Leu Met Val Ser Glu Ser Glu Phe
 1010 1015 1020
 Asp Ser Asp Gln Asp Thr Ile Phe Ser Arg Glu Lys Met Glu Arg Gly
 1025 1030 1035 1040
 Asn Pro Lys Val Ser Met Asn Gly Ser Ile Arg Asn Gly Ala Ser Phe
 1045 1050 1055
 Ser Tyr Cys Ser Lys Asp Arg
 1060

<210> 11
 <211> 1072
 <212> PRT
 <213> Homo sapiens

<400> 11
 Met Ala Pro Pro Thr Gly Val Leu Ser Ser Leu Leu Leu Val Thr
 1 5 10 15
 Ile Ala Gly Cys Ala Arg Lys Gln Cys Ser Glu Gly Arg Thr Tyr Ser
 20 25 30
 Asn Ala Val Ile Ser Pro Asn Leu Glu Thr Thr Arg Ile Met Arg Val
 35 40 45
 Ser His Thr Phe Pro Val Val Asp Cys Thr Ala Ala Cys Cys Asp Leu
 50 55 60
 Ser Ser Cys Asp Leu Ala Trp Trp Phe Glu Gly Arg Cys Tyr Leu Val
 65 70 75 80
 Ser Cys Pro His Lys Glu Asn Cys Glu Pro Lys Lys Met Gly Pro Ile
 85 90 95
 Arg Ser Tyr Leu Thr Phe Val Leu Arg Pro Val Gln Arg Pro Ala Gln
 100 105 110
 Leu Leu Asp Tyr Gly Asp Met Met Leu Asn Arg Gly Ser Pro Ser Gly
 115 120 125
 Ile Trp Gly Asp Ser Pro Glu Asp Ile Arg Lys Asp Leu Thr Phe Leu
 130 135 140
 Gly Lys Asp Trp Gly Leu Glu Glu Met Ser Glu Tyr Ser Asp Asp Tyr
 145 150 155 160
 Arg Glu Leu Glu Lys Asp Leu Leu Gln Pro Ser Gly Lys Gln Glu Pro
 165 170 175
 Arg Gly Ser Ala Glu Tyr Thr Asp Trp Gly Leu Leu Pro Gly Ser Glu
 180 185 190
 Gly Ala Phe Asn Ser Ser Val Gly Asp Ser Pro Ala Val Pro Ala Glu
 195 200 205
 Thr Gln Gln Asp Pro Glu Leu His Tyr Leu Asn Glu Ser Ala Ser Thr
 210 215 220
 Pro Ala Pro Lys Leu Pro Glu Arg Ser Val Leu Leu Pro Leu Pro Thr
 225 230 235 240
 Thr Pro Ser Ser Gly Glu Val Leu Glu Lys Glu Lys Ala Ser Gln Leu
 245 250 255
 Gln Glu Gln Ser Ser Asn Ser Ser Gly Lys Glu Val Leu Met Pro Ser
 260 265 270
 His Ser Leu Pro Pro Ala Ser Leu Glu Leu Ser Ser Val Thr Val Glu
 275 280 285
 Lys Ser Pro Val Leu Thr Val Thr Pro Gly Ser Thr Glu His Ser Ile
 290 295 300
 Pro Thr Pro Pro Thr Ser Ala Ala Pro Ser Glu Ser Thr Pro Ser Glu
 305 310 315 320
 Leu Pro Ile Ser Pro Thr Thr Ala Pro Arg Thr Val Lys Glu Leu Thr
 325 330 335
 Val Ser Ala Gly Asp Asn Leu Ile Ile Thr Leu Pro Asp Asn Glu Val
 340 345 350
 Glu Leu Lys Ala Phe Val Ala Pro Ala Pro Pro Val Glu Thr Thr Tyr
 355 360 365

Asn Tyr Glu Trp Asn Leu Ile Ser His Pro Thr Asp Tyr Gln Gly Glu
 370 375 380
 Ile Lys Gln Gly His Lys Gln Thr Leu Asn Leu Ser Gln Leu Ser Val
 385 390 395 400
 Gly Leu Tyr Val Phe Lys Val Thr Val Ser Ser Glu Asn Ala Phe Gly
 405 410 415
 Glu Gly Phe Val Asn Val Thr Val Lys Pro Ala Arg Arg Val Asn Leu
 420 425 430
 Pro Pro Val Ala Val Val Ser Pro Gln Leu Gln Glu Leu Thr Leu Pro
 435 440 445
 Leu Thr Ser Ala Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr Glu
 450 455 460
 Ile Val Ser Tyr His Trp Glu Glu Ile Asn Gly Pro Phe Ile Glu Glu
 465 470 475 480
 Lys Thr Ser Val Asp Ser Pro Val Leu Arg Leu Ser Asn Leu Asp Pro
 485 490 495
 Gly Asn Tyr Ser Phe Arg Leu Thr Val Thr Asp Ser Asp Gly Ala Thr
 500 505 510
 Asn Ser Thr Thr Ala Ala Leu Ile Val Asn Asn Ala Val Asp Tyr Pro
 515 520 525
 Pro Val Ala Asn Ala Gly Pro Asn His Thr Ile Thr Leu Pro Gln Asn
 530 535 540
 Ser Ile Thr Leu Asn Gly Asn Gln Ser Ser Asp Asp His Gln Ile Val
 545 550 555 560
 Leu Tyr Glu Trp Ser Leu Gly Pro Gly Ser Glu Gly Lys His Val Val
 565 570 575
 Met Gln Gly Val Gln Thr Pro Tyr Leu His Leu Ser Ala Met Gln Glu
 580 585 590
 Gly Asp Tyr Thr Phe Gln Leu Lys Val Thr Asp Ser Ser Arg Gln Gln
 595 600 605
 Ser Thr Ala Val Val Thr Val Ile Val Gln Pro Glu Asn Asn Arg Pro
 610 615 620
 Pro Val Ala Val Ala Gly Pro Asp Lys Glu Leu Ile Phe Pro Val Glu
 625 630 635 640
 Ser Ala Thr Leu Asp Gly Ser Ser Ser Asp Asp His Gly Ile Val
 645 650 655
 Phe Tyr His Trp Glu His Val Arg Gly Pro Ser Ala Val Glu Met Glu
 660 665 670
 Asn Ile Asp Lys Ala Ile Ala Thr Val Thr Gly Leu Gln Val Gly Thr
 675 680 685
 Tyr His Phe Arg Leu Thr Val Lys Asp Gln Gln Gly Leu Ser Ser Thr
 690 695 700
 Ser Thr Leu Thr Val Ala Val Lys Lys Glu Asn Asn Ser Pro Pro Arg
 705 710 715 720
 Ala Arg Ala Gly Gly Arg His Val Leu Val Leu Pro Asn Asn Ser Ile
 725 730 735
 Thr Leu Asp Gly Ser Arg Ser Thr Asp Asp Gln Arg Ile Val Ser Tyr
 740 745 750
 Leu Trp Ile Arg Asp Gly Gln Ser Pro Ala Ala Gly Asp Val Ile Asp
 755 760 765
 Gly Ser Asp His Ser Val Ala Leu Gln Leu Thr Asn Leu Val Glu Gly
 770 775 780
 Val Tyr Thr Phe His Leu Arg Val Thr Asp Ser Gln Gly Ala Ser Asp
 785 790 795 800
 Thr Asp Thr Ala Thr Val Glu Val Gln Pro Asp Pro Arg Lys Ser Gly
 805 810 815
 Leu Val Glu Leu Thr Leu Gln Val Gly Val Gly Gln Leu Thr Glu Gln
 820 825 830
 Arg Lys Asp Thr Leu Val Arg Gln Leu Ala Val Leu Leu Asn Val Leu
 835 840 845
 Asp Ser Asp Ile Lys Val Gln Lys Ile Arg Ala His Ser Asp Leu Ser
 850 855 860
 Thr Val Ile Val Phe Tyr Val Gln Ser Arg Pro Pro Phe Lys Val Leu
 865 870 880

Lys Ala Ala Glu Val Ala Arg Asn Leu His Met Arg Leu Ser Lys Glu
 885 890 895
 Lys Ala Asp Phe Leu Leu Phe Lys Val Leu Arg Val Asp Thr Ala Gly
 900 905 910
 Cys Leu Leu Lys Cys Ser Gly His Gly His Cys Asp Pro Leu Thr Lys
 915 920 925
 Arg Cys Ile Cys Ser His Leu Trp Met Glu Asn Leu Ile Gln Arg Tyr
 930 935 940
 Ile Trp Asp Gly Glu Ser Asn Cys Glu Trp Ser Ile Phe Tyr Val Thr
 945 950 955 960
 Val Leu Ala Phe Thr Leu Ile Val Leu Thr Gly Gly Phe Thr Trp Leu
 965 970 975
 Cys Ile Cys Cys Lys Arg Gln Lys Arg Thr Lys Ile Arg Lys Lys
 980 985 990
 Thr Lys Tyr Thr Ile Leu Asp Asn Met Asp Glu Gln Glu Arg Met Glu
 995 1000 1005
 Leu Arg Pro Lys Tyr Gly Ile Lys His Arg Ser Thr Glu His Asn Ser
 1010 1015 1020
 Ser Leu Met Val Ser Glu Ser Glu Phe Asp Ser Asp Gln Asp Thr Ile
 1025 1030 1035 1040
 Phe Ser Arg Glu Lys Met Glu Arg Gly Asn Pro Lys Val Ser Met Asn
 1045 1050 1055
 Gly Ser Ile Arg Asn Gly Ala Ser Phe Ser Tyr Cys Ser Lys Asp Arg
 1060 1065 1070

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<400> 12

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Asn Ala Val Ile Ser Pro	Asn Leu Glu Thr	Thr Arg Ile	Met Arg Val
35 40	45		
Ser His Thr Phe Pro Val	Val Asp Cys Thr	Ala Ala Cys Cys	Asp Leu
50 55	60		
Ser Ser Cys Asp Leu Ala	Trp Trp Phe Glu	Gly Arg Cys Tyr	Leu Val
65 70	75	80	
Ser Cys Pro His Lys Glu	Asn Cys Glu Pro	Lys Lys Met Gly	Pro Ile
85 90	95		
Arg Ser Tyr Leu Thr Phe	Val Leu Arg Pro	Val Gln Arg Pro	Ala Gln
100 105	110		
Leu Leu Asp Tyr Gly Asp	Met Met Leu Asn	Arg Gly Ser Pro	Ser Gly
115 120	125		
Ile Trp Gly Asp Ser Pro	Glu Asp Ile Arg	Lys Asp Leu Pro	Phe Leu
130 135	140		
Gly Lys Asp Trp Gly Leu	Glu Met Ser Glu	Tyr Ala Asp Asp	Tyr
145 150	155	160	
Arg Glu Leu Glu Lys Asp	Leu Leu Gln Pro	Ser Gly Lys Gln	Glu Pro
165 170	175		
Arg Gly Ser Ala Glu Tyr	Thr Asp Trp Gly	Leu Leu Pro Gly	Ser Glu
180 185	190		
Gly Ala Phe Asn Ser Ser	Val Gly Asp Ser	Pro Ala Val	Pro Ala Glu
195 200	205		
Thr Gln Gln Asp Pro Glu	Leu His Tyr Leu	Asn Glu Ser Ala	Ser Thr
210 215	220		
Pro Ala Pro Lys Leu Pro	Glu Arg Ser Val	Leu Leu Pro	Leu Pro Thr
225 230	235	240	
Thr Pro Ser Ser Gly Glu	Val Leu Glu Lys	Glu Lys Ala Ser	Gln Leu
245 250	255		

Gln Glu Gln Ser Ser Asn Ser Ser Gly Lys Glu Val Leu Met Pro Ser
 260 265 270
 His Ser Leu Pro Pro Ala Ser Leu Glu Leu Ser Ser Val Thr Val Glu
 275 280 285
 Lys Ser Pro Val Leu Thr Val Thr Pro Gly Ser Thr Glu His Ser Ile
 290 295 300
 Pro Thr Pro Pro Thr Ser Ala Ala Pro Ser Glu Ser Thr Pro Ser Glu
 305 310 315 320
 Leu Pro Ile Ser Pro Thr Thr Ala Pro Arg Thr Val Lys Glu Leu Thr
 325 330 335
 Val Ser Ala Gly Asp Asn Leu Ile Ile Thr Leu Pro Asp Asn Glu Val
 340 345 350
 Glu Leu Lys Ala Phe Val Ala Pro Ala Pro Pro Val Glu Thr Thr Tyr
 355 360 365
 Asn Tyr Glu Trp Asn Leu Ile Ser His Pro Thr Asp Tyr Gln Gly Glu
 370 375 380
 Ile Lys Gln Gly His Lys Gln Thr Leu Asn Leu Ser Gln Leu Ser Val
 385 390 395 400
 Gly Leu Tyr Val Phe Lys Val Thr Val Ser Ser Glu Asn Ala Phe Gly
 405 410 415
 Glu Gly Phe Val Asn Val Thr Val Lys Pro Ala Arg Arg Val Asn Leu
 420 425 430
 Pro Pro Val Ala Val Val Ser Pro Gln Leu Gln Glu Leu Thr Leu Pro
 435 440 445
 Leu Thr Ser Ala Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr Glu
 450 455 460
 Ile Val Ser Tyr His Trp Glu Glu Ile Asn Gly Pro Phe Ile Glu Glu
 465 470 475 480
 Lys Thr Ser Val Asp Ser Pro Val Leu Arg Leu Ser Asn Leu Asp Pro
 485 490 495
 Gly Asn Tyr Ser Phe Arg Leu Thr Val Thr Asp Ser Asp Gly Ala Thr
 500 505 510
 Asn Ser Thr Thr Ala Ala Leu Ile Val Asn Asn Ala Val Asp Tyr Pro
 515 520 525
 Pro Val Ala Asn Ala Gly Pro Asn His Thr Ile Thr Leu Pro Gln Asn
 530 535 540
 Ser Ile Thr Leu Asn Gly Asn Gln Ser Ser Asp Asp His Gln Ile Val
 545 550 555 560
 Leu Tyr Glu Trp Ser Leu Gly Pro Gly Ser Glu Gly Lys His Val Val
 565 570 575
 Met Gln Gly Val Gln Thr Pro Tyr Leu His Leu Ser Ala Met Gln Glu
 580 585 590
 Gly Asp Tyr Thr Phe Gln Leu Lys Val Thr Asp Ser Ser Arg Gln Gln
 595 600 605
 Ser Thr Ala Val Val Thr Val Ile Val Gln Pro Glu Asn Asn Arg Pro
 610 615 620
 Pro Val Ala Val Ala Gly Pro Asp Lys Glu Leu Ile Phe Pro Val Glu
 625 630 635 640
 Ser Ala Thr Leu Asp Gly Ser Ser Ser Asp Asp His Gly Ile Val
 645 650 655
 Phe Tyr His Trp Glu His Val Arg Gly Pro Ser Ala Val Glu Met Glu
 660 665 670
 Asn Ile Asp Lys Ala Ile Ala Thr Val Thr Gly Leu Gln Val Gly Thr
 675 680 685
 Tyr His Phe Arg Leu Thr Val Lys Asp Gln Gln Gly Leu Ser Ser Thr
 690 695 700
 Ser Thr Leu Thr Val Ala Val Lys Lys Glu Asn Asn Ser Pro Pro Arg
 705 710 715 720
 Ala Arg Ala Gly Gly Arg His Val Leu Val Leu Pro Asn Asn Ser Ile
 725 730 735
 Thr Leu Asp Gly Ser Arg Ser Thr Asp Asp Gln Arg Ile Val Ser Tyr
 740 745 750
 Leu Trp Ile Arg Asp Gly Gln Ser Pro Ala Ala Gly Asp Val Ile Asp
 755 760 765

Gly	Ser	Asp	His	Ser	Val	Ala	Leu	Gln	Leu	Thr	Asn	Leu	Val	Glu	Gly
770														780	
Val	Tyr	Thr	Phe	His	Leu	Arg	Val	Thr	Asp	Ser	Gln	Gly	Ala	Ser	Asp
785											795				800
Thr	Asp	Thr	Ala	Thr	Val	Glu	Val	Gln	Pro	Asp	Pro	Arg	Lys	Ser	Gly
						805				810				815	
Leu	Val	Glu	Leu	Thr	Leu	Gln	Val	Gly	Val	Gly	Gln	Leu	Thr	Glu	Gln
						820			825				830		
Arg	Lys	Asp	Thr	Leu	Val	Arg	Gln	Leu	Ala	Val	Leu	Leu	Asn	Val	Leu
						835			840			845			
Asp	Ser	Asp	Ile	Lys	Val	Gln	Lys	Ile	Arg	Ala	His	Ser	Asp	Leu	Ser
						850			855			860			
Thr	Val	Ile	Val	Phe	Tyr	Val	Gln	Ser	Arg	Pro	Pro	Phe	Lys	Val	Leu
						865			870			875			880
Lys	Ala	Ala	Glu	Val	Ala	Arg	Asn	Leu	His	Met	Arg	Leu	Ser	Lys	Glu
						885			890			895			
Lys	Ala	Asp	Phe	Leu	Leu	Phe	Lys	Val	Leu	Arg	Val	Asp	Thr	Ala	Gly
						900			905			910			
Cys	Leu	Leu	Lys	Cys	Ser	Gly	His	Gly	His	Cys	Asp	Pro	Leu	Thr	Lys
						915			920			925			
Arg	Cys	Ile	Cys	Ser	His	Leu	Trp	Met	Glu	Asn	Leu	Ile	Gln	Arg	Tyr
						930			935			940			
Ile	Trp	Asp	Gly	Glu	Ser	Asn	Cys	Glu	Trp	Ser	Ile	Phe	Tyr	Val	Thr
						945			950			955			960
Val	Leu	Ala	Phe	Thr	Leu	Ile	Val	Leu	Thr	Gly	Gly	Phe	Thr	Trp	Leu
						965			970			975			
Cys	Ile	Cys	Cys	Lys	Arg	Gln	Lys	Arg	Thr	Lys	Ile	Arg	Lys	Lys	
						980			985			990			
Thr	Lys	Tyr	Thr	Ile	Leu	Asp	Asn	Met	Asp	Glu	Gln	Glu	Arg	Met	Glu
						995			1000			1005			
Leu	Arg	Pro	Lys	Tyr	Gly	Ile	Lys	His	Arg	Ser	Thr	Glu	His	Asn	Ser
						1010			1015			1020			
Ser	Leu	Met	Val	Ser	Glu	Ser	Glu	Phe	Asp	Ser	Asp	Gln	Asp	Thr	Ile
						1025			1030			1035			1040
Phe	Ser	Arg	Glu	Lys	Met	Glu	Arg	Gly	Asn	Pro	Lys	Val	Ser	Met	Asn
						1045			1050			1055			
Gly	Ser	Ile	Arg	Asn	Gly	Ala	Ser	Phe	Ser	Tyr	Cys	Ser	Lys	Asp	Arg
						1060			1065			1070			

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<211> 21
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Asn Val Val Asn Ser
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<210> 15
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 <223> Xaa = D-alanine or L-alanine

<221> VARIANT
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<400> 16
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 <213> Artificial Sequence

<220>
 <223> cDNA synthesis primer

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<210> 18
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<220>
 <223> Primer

<400> 18
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<220>
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<400> 19
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<210> 20

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<220>
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gatcctcggc 10

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<400> 47
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<210> 54
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<400> 54
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Ala Phe Asn Ser Ser Val Gly Asp Ser
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Tyr Leu Asn Glu Ser Ala Ser Thr Pro
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Thr Thr Pro Ser Ser Gly Glu Val Leu
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<210> 73
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<400> 73
Met Pro Ser His Ser Leu Pro Pro Ala
1 5

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<400> 74
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1 5

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Leu Glu Leu Ser Ser Val Thr Val Glu
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Thr Val Glu Lys Ser Pro Val Leu Thr
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Val Thr Pro Gly Ser Thr Glu His Ser
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Glu Ser Thr Pro Ser Glu Leu Pro Ile
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Glu Leu Pro Ile Ser Pro Thr Thr Ala
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Glu Leu Thr Val Ser Ala Gly Asp Asn
1 5

<210> 86

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Trp Asn Leu Ile Ser His Pro Thr Asp
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<210> 87

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<212> PRT

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<400> 87

Thr Leu Asn Leu Ser Gln Leu Ser Val
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<210> 88

<211> 9

<212> PRT

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<400> 88

Leu Ser Gln Leu Ser Val Gly Leu Tyr
1 5

<210> 89

<211> 9

<212> PRT

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Lys Val Thr Val Ser Ser Glu Asn Ala

<210> 90
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<400> 90
Val Thr Val Ser Ser Glu Asn Ala Phe
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Val Ala Val Val Ser Pro Gln Leu Gln
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<210> 92
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Leu Pro Leu Thr Ser Ala Leu Ile Asp
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<213> Homo sapiens

<400> 250
Gln Val Gly Thr Tyr His Phe Arg Leu
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<400> 251
Arg Ile Val Ser Tyr Leu Trp Ile Arg
1 5

<210> 252
<211> 9
<212> PRT
<213> Homo sapiens

<400> 252
Val Glu Gly Val Tyr Thr Phe His Leu
1 5

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<400> 254
Leu Ile Gln Arg Tyr Ile Trp Asp Gly
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<213> Homo sapiens

<400> 255
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<400> 256
 Lys Lys Thr Lys Tyr Thr Ile Leu Asp
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<400> 257
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 <212> PRT
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 35 40 45
 Ser His Thr Phe Pro Val Val Asp Cys Thr Ala Ala Cys Cys Asp Leu
 50 55 60
 Ser Ser Cys Asp Leu Ala Trp Trp Phe Glu Gly Arg Cys Tyr Leu Val
 65 70 75 80
 Ser Cys Pro His Lys Glu Asn Cys Glu Pro Lys Lys Met Gly Pro Ile
 85 90 95
 Arg Ser Tyr Leu Thr Phe Val Leu Arg Pro Val Gln Arg Pro Ala Gln
 100 105 110
 Leu Leu Asp Tyr Gly Asp Met Met Leu Asn Arg Gly Ser Pro Ser Gly
 115 120 125
 Ile Trp Gly Asp Ser Pro Glu Asp Ile Arg Lys Asp Leu Pro Phe Leu
 130 135 140
 Gly Lys Asp Trp Gly Leu Glu Met Ser Glu Tyr Ser Asp Asp Tyr
 145 150 155 160
 Arg Glu Leu Glu Lys Asp Leu Leu Gln Pro Ser Gly Lys Gln Glu Pro
 165 170 175
 Arg Gly Ser Ala Glu Tyr Thr Asp Trp Gly Leu Leu Pro Gly Ser Glu
 180 185 190
 Gly Ala Phe Asn Ser Ser Val Gly Asp Ser Pro Ala Val Pro Ala Glu
 195 200 205
 Thr Gln Gln Asp Pro Glu Leu His Tyr Leu Asn Glu Ser Ala Ser Thr
 210 215 220
 Pro Ala Pro Lys Leu Pro Glu Arg Ser Val Leu Leu Pro Leu Pro Thr

225	230	235	240
Thr Pro Ser Ser Gly Glu Val Leu Glu Lys	Glu Lys Ala Ser Gln Leu		
245	250		255
Gln Glu Gln Ser Ser Asn Ser Ser Gly Lys	Glu Val Leu Met Pro Ser		
260	265	270	
His Ser Leu Pro Pro Ala Ser Leu Glu Leu Ser Ser Val	Thr Val Glu		
275	280	285	
Lys Ser Pro Val Leu Thr Val Thr Pro Gly Ser	Thr Glu His Ser Ile		
290	295	300	
Pro Thr Pro Pro Thr Ser Ala Ala Pro Ser Glu	Ser Thr Pro Ser Glu		
305	310	315	320
Leu Pro Ile Ser Pro Thr Thr Ala Pro Arg	Thr Val Lys Glu Leu Thr		
325	330		335
Val Ser Ala Gly Asp Asn Leu Ile Ile Thr	Leu Pro Asp Asn Glu Val		
340	345	350	
Glu Leu Lys Ala Phe Val Ala Pro Ala Pro Val	Glu Thr Thr Tyr		
355	360	365	
Asn Tyr Glu Trp Asn Leu Ile Ser His Pro Thr	Asp Tyr Gln Gly Glu		
370	375	380	
Ile Lys Gln Gly His Lys Gln Thr Leu Asn	Leu Ser Gln Leu Ser Val		
385	390	395	400
Gly Leu Tyr Val Phe Lys Val Thr Val Ser	Ser Glu Asn Ala Phe Gly		
405	410		415
Glu Gly Phe Val Asn Val Thr Val Lys Pro Ala Arg	Arg Val Asn Leu		
420	425	430	
Pro Pro Val Ala Val Val Ser Pro Gln Leu Gln	Glu Leu Thr Leu Pro		
435	440	445	
Leu Thr Ser Ala Leu Ile Asp Gly Ser Gln Ser	Thr Asp Asp Thr Glu		
450	455	460	
Ile Val Ser Tyr His Trp Glu Glu Ile Asn Gly	Pro Phe Ile Glu Glu		
465	470	475	480
Lys Thr Ser Val Asp Ser Pro Val Leu Arg	Leu Ser Asn Leu Asp Pro		
485	490	495	
Gly Asn Tyr Ser Phe Arg Leu Thr Val Thr Asp	Ser Asp Gly Ala Thr		
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Asn Ser Thr Thr Ala Ala Leu Ile Val Asn Asn	Ala Val Asp Tyr Pro		
515	520	525	
Pro Val Ala Asn Ala Gly Pro Asn His Thr Ile	Thr Leu Pro Gln Asn		
530	535	540	
Ser Ile Thr Leu Asn Gly Asn Gln Ser Ser Asp	Asp His Gln Ile Val		
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Leu Tyr Glu Trp Ser Leu Gly Pro Gly Ser	Glu Gly Lys His Val Val		
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Met Gln Gly Val Gln Thr Pro Tyr Leu His	Leu Ser Ala Met Gln Glu		
580	585	590	
Gly Asp Tyr Thr Phe Gln Leu Lys Val Thr Asp	Ser Ser Arg Gln Gln		
595	600	605	
Ser Thr Ala Val Val Thr Val Ile Val Gln Pro	Glu Asn Asn Arg Pro		
610	615	620	
Pro Val Ala Val Ala Gly Pro Asp Lys Glu	Leu Ile Phe Pro Val Glu		
625	630	635	640
Ser Ala Thr Leu Asp Gly Ser Ser Ser Asp	Asp His Gly Ile Val		
645	650	655	
Phe Tyr His Trp Glu His Val Arg Gly	Pro Ser Ala Val Glu Met Glu		
660	665	670	
Asn Ile Asp Lys Ala Ile Ala Thr Val Thr	Gly Leu Gln Val Gly Thr		
675	680	685	
Tyr His Phe Arg Leu Thr Val Lys Asp Gln Gln	Gly Leu Ser Ser Thr		
690	695	700	
Ser Thr Leu Thr Val Ala Val Lys Lys Glu	Asn Asn Ser Pro Pro Arg		
705	710	715	720
Ala Arg Ala Gly Gly Arg His Val Leu Val Leu	Pro Asn Asn Ser Ile		
725	730	735	
Thr Leu Asp Gly Ser Arg Ser Thr Asp Asp Gln	Arg Ile Val Ser Tyr		

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Leu Trp Ile Arg Asp Gly Gln Ser Pro Ala Ala Gly Asp Val Ile Asp	755	760	765
Gly Ser Asp His Ser Val Ala Leu Gln Leu Thr Asn Leu Val Glu Gly	770	775	780
Val Tyr Thr Phe His Leu Arg Val Thr Asp Ser Gln Gly Ala Ser Asp	785	790	795
Thr Asp Thr Ala Thr Val Glu Val Gln Pro Asp Pro Arg Lys Ser Gly	805	810	815
Leu Val Glu Leu Thr Leu Gln Val Gly Val Gly Gln Leu Thr Glu Gln	820	825	830
Arg Lys Asp Thr Leu Val Arg Gln Leu Ala Val Leu Leu Asn Val Leu	835	840	845
Asp Ser Asp Ile Lys Val Gln Lys Ile Arg Ala His Ser Asp Leu Ser	850	855	860
Thr Val Ile Val Phe Tyr Val Gln Ser Arg Pro Pro Phe Lys Val Leu	865	870	875
Lys Ala Ala Glu Val Ala Arg Asn Leu His Met Arg Leu Ser Lys Glu	885	890	895
Lys Ala Asp Phe Leu Leu Phe Lys Val Leu Arg Val Asp Thr Ala Gly	900	905	910
Cys Leu Leu Lys Cys Ser Gly His Gly His Cys Asp Pro Leu Thr Lys	915	920	925
Arg Cys Ile Cys Ser His Leu Trp Met Glu Asn Leu Ile Gln Arg Tyr	930	935	940
Ile Trp Asp Gly Glu Ser Asn Cys Glu Trp Ser Ile Phe Tyr Val Thr	945	950	955
Val Leu Ala Phe Thr Leu Ile Val Leu Thr Gly Gly Phe Thr Trp Leu	965	970	975
Cys Ile Cys Cys Cys Lys Arg Gln Lys Arg Thr Lys Ile Arg Lys Lys	980	985	990
Thr Lys Tyr Thr Ile Leu Asp Asn Met Asp Glu Gln Glu Arg Met Glu	995	1000	1005
Leu Arg Pro Lys Tyr Gly Ile Lys His Arg Ser Thr Glu His Asn Ser	1010	1015	1020
Ser Leu Met Val Ser Glu Ser Glu Phe Asp Ser Asp Gln Asp Thr Ile	1025	1030	1035
Phe Ser Arg Glu Lys Met Glu Arg Gly Asn Pro Lys Val Ser Met Asn	1045	1050	1055
Gly Ser Ile Arg Asn Gly Ala Ser Phe Ser Tyr Cys Ser Lys Asp Arg	1060	1065	1070

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<211> 17

<212> PRT

<213> Homo sapiens

<400> 260

Gly Leu Glu Glu Met Ser Glu Tyr Ala Asp Asp Tyr Arg Glu Leu Glu	1	5	10	15
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Lys

<210> 261

<211> 19

<212> PRT

<213> Homo sapiens

<400> 261

Trp Gly Leu Glu Glu Met Ser Glu Tyr Ala Asp Asp Tyr Arg Glu Leu	1	5	10	15
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Glu Lys Asp

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<211> 29
<212> PRT
<213> Homo sapiens

<400> 262
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Asp Tyr Arg Glu Leu Glu Lys Asp Leu Leu Gln Pro Ser
20 25

<210> 263
<211> 18
<212> PRT
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<400> 263
Met Thr Arg Leu Gly Trp Pro Ser Pro Cys Cys Ala Arg Lys Gln Cys
1 5 10 15
Ser Glu

<210> 264
<211> 19
<212> PRT
<213> Homo sapiens

<400> 264
Met Thr Arg Leu Gly Trp Pro Ser Pro Cys Cys Ala Arg Lys Gln Cys
1 5 10 15
Ser Glu Gly

<210> 265
<211> 24
<212> PRT
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<400> 265
Met Thr Arg Leu Gly Trp Pro Ser Pro Cys Cys Ala Arg Lys Gln Cys
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Ser Glu Gly Arg Thr Tyr Ser Asn
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<210> 266
<211> 17
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<400> 266
Pro Glu Asp Ile Arg Lys Asp Leu Thr Phe Leu Gly Lys Asp Trp Gly
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<210> 267

<211> 19
 <212> PRT
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<400> 267
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 1 5 10 15
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<210> 268
 <211> 29
 <212> PRT
 <213> Homo sapiens

<400> 268
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 <212> DNA
 <213> Homo sapiens

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Ser	Glu	Gly	Lys	His	Val	Val	Met	Gln	Gly	Val	Gln	Thr	Pro	Tyr	Leu
565															
His	Leu	Ser	Ala	Met	Gln	Glu	Gly	Asp	Tyr	Thr	Phe	Gln	Leu	Lys	Val
580															
Thr	Asp	Ser	Ser	Arg	Gln	Gln	Ser	Thr	Ala	Val	Val	Thr	Val	Ile	Val
595															
Gln	Pro	Glu	Asn	Asn	Arg	Pro	Pro	Val	Ala	Val	Ala	Gly	Pro	Asp	Lys
610															
Glu	Leu	Ile	Phe	Pro	Val	Glu	Ser	Ala	Thr	Leu	Asp	Gly	Ser	Ser	Ser
625															
Ser	Asp	Asp	His	Gly	Ile	Val	Phe	Tyr	His	Trp	Glu	His	Val	Arg	Gly
645															
Pro	Ser	Ala	Val	Glu	Met	Glu	Asn	Ile	Asp	Lys	Ala	Ile	Ala	Thr	Val
660															
Thr	Gly	Leu	Gln	Val	Gly	Thr	Tyr	His	Phe	Arg	Leu	Thr	Val	Lys	Asp
675															
Gln	Gln	Gly	Leu	Ser	Ser	Thr	Ser	Thr	Leu	Thr	Val	Ala	Val	Lys	Lys
690															
Glu	Asn	Asn	Ser	Pro	Pro	Arg	Ala	Arg	Ala	Gly	Gly	Arg	His	Val	Leu
705															
Val	Leu	Pro	Asn	Asn	Ser	Ile	Thr	Leu	Asp	Gly	Ser	Arg	Ser	Thr	Asp
725															
Asp	Gln	Arg	Ile	Val	Ser	Tyr	Leu	Trp	Ile	Arg	Asp	Gly	Gln	Ser	Pro
740															
Ala	Ala	Gly	Asp	Val	Ile	Asp	Gly	Ser	Asp	His	Ser	Val	Ala	Gln	
755															
Leu	Thr	Asn	Leu	Val	Glu	Gly	Val	Tyr	Thr	Phe	His	Leu	Arg	Val	Thr
770															
Asp	Ser	Gln	Gly	Ala	Ser	Asp	Thr	Asp	Thr	Ala	Thr	Val	Glu	Val	Gln
785															
Pro	Asp	Pro	Arg	Lys	Ser	Gly	Leu	Val	Glu	Leu	Thr	Leu	Gln	Val	Gly
805															
Val	Gly	Gln	Leu	Thr	Glu	Gln	Arg	Lys	Asp	Thr	Leu	Val	Arg	Gln	Leu
820															
Ala	Val	Leu	Leu	Asn	Val	Leu	Asp	Ser	Asp	Ile	Lys	Val	Gln	Lys	Ile
835															
Arg	Ala	His	Ser	Asp	Leu	Ser	Thr	Val	Ile	Val	Phe	Tyr	Val	Gln	Ser
850															
Arg	Pro	Pro	Phe	Lys	Val	Leu	Lys	Ala	Ala	Glu	Val	Ala	Arg	Asn	Leu
865															
His	Met	Arg	Leu	Ser	Lys	Glu	Lys	Ala	Asp	Phe	Leu	Leu	Phe	Lys	Val

885	890	895
Leu Arg Val Asp Thr Ala Gly Cys Leu	Leu Lys Cys Ser Gly His Gly	
900	905	910
His Cys Asp Pro Leu Thr Lys Arg Cys Ile Cys Ser His	Leu Trp Met	
915	920	925
Glu Asn Leu Ile Gln Arg Tyr Ile Trp Asp Gly Glu	Ser Asn Cys Glu	
930	935	940
Trp Ser Ile Phe Tyr Val Thr Val Leu Ala Phe	Thr Leu Ile Val Leu	
945	950	955
Thr Gly Phe Thr Trp Leu Cys Ile Cys Cys Lys Arg Gln Lys		
965	970	975
Arg Thr Lys Ile Arg Lys Lys Thr Lys Tyr Thr Ile Leu Asp Asn Met		
980	985	990
Asp Glu Gln Glu Arg Met Glu Leu Arg Pro Lys Tyr Gly Ile Lys His		
995	1000	1005
Arg Ser Thr Glu His Asn Ser Ser Leu Met Val Ser Glu Ser Glu Phe		
1010	1015	1020
Asp Ser Asp Gln Asp Thr Ile Phe Ser Arg Glu Lys Met Glu Arg Gly		
1025	1030	1035
Asn Pro Lys Val Ser Met Asn Gly Ser Ile Arg Asn Gly Ala Ser Phe		
1045	1050	1055
Ser Tyr Cys Ser Lys Asp Arg		
1060		

<210> 273
<211> 1053
<212> PRT
<213> Homo sapiens

<400> 273		
Cys Ala Arg Lys Gln Cys Ser Glu Gly Arg Thr Tyr Ser Asn Ala Val		
1	5	10
Ile Ser Pro Asn Leu Glu Thr Thr Arg Ile Met Arg Val Ser His Thr		
20	25	30
Phe Pro Val Val Asp Cys Thr Ala Ala Cys Cys Asp Leu Ser Ser Cys		
35	40	45
Asp Leu Ala Trp Trp Phe Glu Gly Arg Cys Tyr Leu Val Ser Cys Pro		
50	55	60
His Lys Glu Asn Cys Glu Pro Lys Lys Met Gly Pro Ile Arg Ser Tyr		
65	70	75
Leu Thr Phe Val Leu Arg Pro Val Gln Arg Pro Ala Gln Leu Leu Asp		
85	90	95
Tyr Gly Asp Met Met Leu Asn Arg Gly Ser Pro Ser Gly Ile Trp Gly		
100	105	110
Asp Ser Pro Glu Asp Ile Arg Lys Asp Leu Pro Phe Leu Gly Lys Asp		
115	120	125
Trp Gly Leu Glu Glu Met Ser Glu Tyr Ser Asp Asp Tyr Arg Glu Leu		
130	135	140
Glu Lys Asp Leu Leu Gln Pro Ser Gly Lys Gln Glu Pro Arg Gly Ser		
145	150	155
Ala Glu Tyr Thr Asp Trp Gly Leu Leu Pro Gly Ser Glu Gly Ala Phe		
165	170	175
Asn Ser Ser Val Gly Asp Ser Pro Ala Val Pro Ala Glu Thr Gln Gln		
180	185	190
Asp Pro Glu Leu His Tyr Leu Asn Glu Ser Ala Ser Thr Pro Ala Pro		
195	200	205
Lys Leu Pro Glu Arg Ser Val Leu Leu Pro Leu Pro Thr Thr Pro Ser		
210	215	220
Ser Gly Glu Val Leu Glu Lys Glu Lys Ala Ser Gln Leu Gln Glu Gln		
225	230	235
Ser Ser Asn Ser Ser Gly Lys Glu Val Leu Met Pro Ser His Ser Leu		
245	250	255
Pro Pro Ala Ser Leu Glu Leu Ser Ser Val Thr Val Glu Lys Ser Pro		

260	265	270
Val Leu Thr Val Thr Pro Gly Ser	Thr Glu His Ser Ile Pro Thr Pro	
275	280	285
Pro Thr Ser Ala Ala Pro Ser Glu Ser Thr Pro Ser	Glu Leu Pro Ile	
290	295	300
Ser Pro Thr Thr Ala Pro Arg Thr Val Lys Glu	Leu Thr Val Ser Ala	
305	310	315
Gly Asp Asn Leu Ile Ile Thr Leu Pro Asp Asn Glu Val	Glu Leu Lys	
325	330	335
Ala Phe Val Ala Pro Ala Pro Val Glu Thr Thr Tyr	Asn Tyr Glu	
340	345	350
Trp Asn Leu Ile Ser His Pro Thr Asp Tyr Gln Gly	Glu Ile Lys Gln	
355	360	365
Gly His Lys Gln Thr Leu Asn Leu Ser Gln Leu Ser	Val Gly Leu Tyr	
370	375	380
Val Phe Lys Val Thr Val Ser Ser Glu Asn Ala Phe	Gly Glu Gly Phe	
385	390	395
Val Asn Val Thr Val Lys Pro Ala Arg Arg Val Asn	Leu Pro Pro Val	
405	410	415
Ala Val Val Ser Pro Gln Leu Gln Glu Leu Thr Leu	Pro Leu Thr Ser	
420	425	430
Ala Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr	Glu Ile Val Ser	
435	440	445
Tyr His Trp Glu Glu Ile Asn Gly Pro Phe Ile	Glu Glu Lys Thr Ser	
450	455	460
Val Asp Ser Pro Val Leu Arg Leu Ser Asn Leu Asp	Pro Gly Asn Tyr	
465	470	475
Ser Phe Arg Leu Thr Val Thr Asp Ser Asp Gly Ala	Thr Asn Ser Thr	
485	490	495
Thr Ala Ala Leu Ile Val Asn Asn Ala Val Asp	Tyr Pro Pro Val Ala	
500	505	510
Asn Ala Gly Pro Asn His Thr Ile Thr Leu Pro	Gln Asn Ser Ile Thr	
515	520	525
Leu Asn Gly Asn Gln Ser Ser Asp Asp His Gln	Ile Val Leu Tyr Glu	
530	535	540
Trp Ser Leu Gly Pro Gly Ser Glu Gly Lys His	Val Val Met Gln Gly	
545	550	555
Val Gln Thr Pro Tyr Leu His Leu Ser Ala Met	Gln Glu Gly Asp Tyr	
565	570	575
Thr Phe Gln Leu Lys Val Thr Asp Ser Ser Arg	Gln Gln Ser Thr Ala	
580	585	590
Val Val Thr Val Ile Val Gln Pro Glu Asn Asn	Arg Pro Pro Val Ala	
595	600	605
Val Ala Gly Pro Asp Lys Glu Leu Ile Phe Pro	Val Glu Ser Ala Thr	
610	615	620
Leu Asp Gly Ser Ser Ser Asp Asp His	Gly Ile Val Phe Tyr His	
625	630	635
Trp Glu His Val Arg Gly Pro Ser Ala Val Glu	Met Glu Asn Ile Asp	
645	650	655
Lys Ala Ile Ala Thr Val Thr Gly Leu Gln Val	Gly Thr Tyr His Phe	
660	665	670
Arg Leu Thr Val Lys Asp Gln Gln Gly Leu Ser	Ser Thr Ser Thr Leu	
675	680	685
Thr Val Ala Val Lys Lys Glu Asn Asn Ser	Pro Pro Arg Ala Arg Ala	
690	695	700
Gly Gly Arg His Val Leu Val Leu Pro Asn Asn	Ser Ile Thr Leu Asp	
705	710	715
Gly Ser Arg Ser Thr Asp Asp Gln Arg Ile Val	Ser Tyr Leu Trp Ile	
725	730	735
Arg Asp Gly Gln Ser Pro Ala Ala Gly Asp Val	Ile Asp Gly Ser Asp	
740	745	750
His Ser Val Ala Leu Gln Leu Thr Asn Leu Val	Glu Gly Val Tyr Thr	
755	760	765
Phe His Leu Arg Val Thr Asp Ser Gln Gly Ala	Ser Asp Thr Asp Thr	

770	775	780
Ala	Thr Val Glu Val Gln Pro Asp Pro Arg Lys Ser Gly Leu Val Glu	
785	790	795
Leu	Thr Leu Gln Val Gly Val Gly Gln Leu Thr Glu Gln Arg Lys Asp	800
	805	810
Thr	Leu Val Arg Gln Leu Ala Val Leu Leu Asn Val Leu Asp Ser Asp	
	820	825
Ile	Lys Val Gln Lys Ile Arg Ala His Ser Asp Leu Ser Thr Val Ile	
	835	840
Val	Phe Tyr Val Gln Ser Arg Pro Pro Phe Lys Val Leu Lys Ala Ala	
	850	855
Glu	Val Ala Arg Asn Leu His Met Arg Leu Ser Lys Glu Lys Ala Asp	
	865	870
Phe	Leu Leu Phe Lys Val Leu Arg Val Asp Thr Ala Gly Cys Leu Leu	
	885	890
Lys	Cys Ser Gly His Gly His Cys Asp Pro Leu Thr Lys Arg Cys Ile	
	900	905
Cys	Ser His Leu Trp Met Glu Asn Leu Ile Gln Arg Tyr Ile Trp Asp	
	915	920
Gly	Glu Ser Asn Cys Glu Trp Ser Ile Phe Tyr Val Thr Val Leu Ala	
	930	935
Phe	Thr Leu Ile Val Leu Thr Gly Gly Phe Thr Trp Leu Cys Ile Cys	
	945	950
Cys	Cys Lys Arg Gln Lys Arg Thr Lys Ile Arg Lys Lys Thr Lys Tyr	
	965	970
Thr	Ile Leu Asp Asn Met Asp Glu Gln Glu Arg Met Glu Leu Arg Pro	
	980	985
Lys	Tyr Gly Ile Lys His Arg Ser Thr Glu His Asn Ser Ser Leu Met	
	995	1000
Val	Ser Glu Ser Glu Phe Asp Ser Asp Gln Asp Thr Ile Phe Ser Arg	
	1010	1015
Glu	Lys Met Glu Arg Gly Asn Pro Lys Val Ser Met Asn Gly Ser Ile	
	1025	1030
Arg	Asn Gly Ala Ser Phe Ser Tyr Cys Ser Lys Asp Arg	
	1045	1050

<210> 274
 <211> 1053
 <212> PRT
 <213> Homo sapiens

<400> 274
 Cys Ala Arg Lys Gln Cys Ser Glu Gly Arg Thr Tyr Ser Asn Ala Val
 1 5 10 15
 Ile Ser Pro Asn Leu Glu Thr Thr Arg Ile Met Arg Val Ser His Thr
 20 25 30
 Phe Pro Val Val Asp Cys Thr Ala Ala Cys Cys Asp Leu Ser Ser Cys
 35 40 45
 Asp Leu Ala Trp Trp Phe Glu Gly Arg Cys Tyr Leu Val Ser Cys Pro
 50 55 60
 His Lys Glu Asn Cys Glu Pro Lys Lys Met Gly Pro Ile Arg Ser Tyr
 65 70 75 80
 Leu Thr Phe Val Leu Arg Pro Val Gln Arg Pro Ala Gln Leu Leu Asp
 85 90 95
 Tyr Gly Asp Met Met Leu Asn Arg Gly Ser Pro Ser Gly Ile Trp Gly
 100 105 110
 Asp Ser Pro Glu Asp Ile Arg Lys Asp Leu Pro Phe Leu Gly Lys Asp
 115 120 125
 Trp Gly Leu Glu Glu Met Ser Glu Tyr Ser Asp Asp Tyr Arg Glu Leu
 130 135 140
 Glu Lys Asp Leu Leu Gln Pro Ser Gly Lys Gln Glu Pro Arg Gly Ser
 145 150 155 160
 Ala Glu Tyr Thr Asp Trp Gly Leu Leu Pro Gly Ser Glu Gly Ala Phe

165	170	175
Asn Ser Ser Val Gly Asp Ser Pro Ala Val Pro Ala Glu Thr Gln Gln		
180	185	190
Asp Pro Glu Leu His Tyr Leu Asn Glu Ser Ala Ser Thr Pro Ala Pro		
195	200	205
Lys Leu Pro Glu Arg Ser Val Leu Leu Pro Leu Pro Thr Thr Pro Ser		
210	215	220
Ser Gly Glu Val Leu Glu Lys Glu Lys Ala Ser Gln Leu Gln Glu Gln		
225	230	235
Ser Ser Asn Ser Ser Gly Lys Glu Val Leu Met Pro Ser His Ser Leu		
245	250	255
Pro Pro Ala Ser Leu Glu Leu Ser Ser Val Thr Val Glu Lys Ser Pro		
260	265	270
Val Leu Thr Val Thr Pro Gly Ser Thr Glu His Ser Ile Pro Thr Pro		
275	280	285
Pro Thr Ser Ala Ala Pro Ser Glu Ser Thr Pro Ser Glu Leu Pro Ile		
290	295	300
Ser Pro Thr Thr Ala Pro Arg Thr Val Lys Glu Leu Thr Val Ser Ala		
305	310	315
Gly Asp Asn Leu Ile Ile Thr Leu Pro Asp Asn Glu Val Glu Leu Lys		
325	330	335
Ala Phe Val Ala Pro Ala Pro Pro Val Glu Thr Thr Tyr Asn Tyr Glu		
340	345	350
Trp Asn Leu Ile Ser His Pro Thr Asp Tyr Gln Gly Glu Ile Lys Gln		
355	360	365
Gly His Lys Gln Thr Leu Asn Leu Ser Gln Leu Ser Val Gly Leu Tyr		
370	375	380
Val Phe Lys Val Thr Val Ser Ser Glu Asn Ala Phe Gly Glu Gly Phe		
385	390	395
400		
Val Asn Val Thr Val Lys Pro Ala Arg Arg Val Asn Leu Pro Pro Val		
405	410	415
Ala Val Val Ser Pro Gln Leu Gln Glu Leu Thr Leu Pro Leu Thr Ser		
420	425	430
Ala Leu Ile Asp Gly Ser Gln Ser Thr Asp Asp Thr Glu Ile Val Ser		
435	440	445
Tyr His Trp Glu Glu Ile Asn Gly Pro Phe Ile Glu Glu Lys Thr Ser		
450	455	460
Val Asp Ser Pro Val Leu Arg Leu Ser Asn Leu Asp Pro Gly Asn Tyr		
465	470	475
480		
Ser Phe Arg Leu Thr Val Thr Asp Ser Asp Gly Ala Thr Asn Ser Thr		
485	490	495
Thr Ala Ala Leu Ile Val Asn Asn Ala Val Asp Tyr Pro Pro Val Ala		
500	505	510
Asn Ala Gly Pro Asn His Thr Ile Thr Leu Pro Gln Asn Ser Ile Thr		
515	520	525
Leu Asn Gly Asn Gln Ser Ser Asp Asp His Gln Ile Val Leu Tyr Glu		
530	535	540
Trp Ser Leu Gly Pro Gly Ser Glu Gly Lys His Val Val Met Gln Gly		
545	550	555
560		
Val Gln Thr Pro Tyr Leu His Leu Ser Ala Met Gln Glu Gly Asp Tyr		
565	570	575
Thr Phe Gln Leu Lys Val Thr Asp Ser Ser Arg Gln Gln Ser Thr Ala		
580	585	590
Val Val Thr Val Ile Val Gln Pro Glu Asn Asn Arg Pro Pro Val Ala		
595	600	605
Val Ala Gly Pro Asp Lys Glu Leu Ile Phe Pro Val Glu Ser Ala Thr		
610	615	620
Leu Asp Gly Ser Ser Ser Asp Asp His Gly Ile Val Phe Tyr His		
625	630	635
640		
Trp Glu His Val Arg Gly Pro Ser Ala Val Glu Met Glu Asn Ile Asp		
645	650	655
Lys Ala Ile Ala Thr Val Thr Gly Leu Gln Val Gly Thr Tyr His Phe		
660	665	670
Arg Leu Thr Val Lys Asp Gln Gln Gly Leu Ser Ser Thr Ser Thr Leu		

675	680	685
Thr Val Ala Val Lys Lys Glu Asn Asn Ser Pro Pro Arg Ala Arg Ala		
690	695	700
Gly Gly Arg His Val Leu Val Leu Pro Asn Asn Ser Ile Thr Leu Asp		
705	710	715
Gly Ser Arg Ser Thr Asp Asp Gln Arg Ile Val Ser Tyr Leu Trp Ile		720
725	730	735
Arg Asp Gly Gln Ser Pro Ala Ala Gly Asp Val Ile Asp Gly Ser Asp		
740	745	750
His Ser Val Ala Leu Gln Leu Thr Asn Leu Val Glu Gly Val Tyr Thr		
755	760	765
Phe His Leu Arg Val Thr Asp Ser Gln Gly Ala Ser Asp Thr Asp Thr		
770	775	780
Ala Thr Val Glu Val Gln Pro Asp Pro Arg Lys Ser Gly Leu Val Glu		
785	790	795
Leu Thr Leu Gln Val Gly Val Gly Gln Leu Thr Glu Gln Arg Lys Asp		800
805	810	815
Thr Leu Val Arg Gln Leu Ala Val Leu Leu Asn Val Leu Asp Ser Asp		
820	825	830
Ile Lys Val Gln Lys Ile Arg Ala His Ser Asp Leu Ser Thr Val Ile		
835	840	845
Val Phe Tyr Val Gln Ser Arg Pro Pro Phe Lys Val Leu Lys Ala Ala		
850	855	860
Glu Val Ala Arg Asn Leu His Met Arg Leu Ser Lys Glu Lys Ala Asp		
865	870	875
Phe Leu Leu Phe Lys Val Leu Arg Val Asp Thr Ala Gly Cys Leu Leu		
885	890	895
Lys Cys Ser Gly His Gly His Cys Asp Pro Leu Thr Lys Arg Cys Ile		
900	905	910
Cys Ser His Leu Trp Met Glu Asn Leu Ile Gln Arg Tyr Ile Trp Asp		
915	920	925
Gly Glu Ser Asn Cys Glu Trp Ser Ile Phe Tyr Val Thr Val Leu Ala		
930	935	940
Phe Thr Leu Ile Val Leu Thr Gly Gly Phe Thr Trp Leu Cys Ile Cys		
945	950	955
Cys Cys Lys Arg Gln Lys Arg Thr Lys Ile Arg Lys Lys Thr Lys Tyr		
965	970	975
Thr Ile Leu Asp Asn Met Asp Glu Gln Glu Arg Met Glu Leu Arg Pro		
980	985	990
Lys Tyr Gly Ile Lys His Arg Ser Thr Glu His Asn Ser Ser Leu Met		
995	1000	1005
Val Ser Glu Ser Glu Phe Asp Ser Asp Gln Asp Thr Ile Phe Ser Arg		
1010	1015	1020
Glu Lys Met Glu Arg Gly Asn Pro Lys Val Ser Met Asn Gly Ser Ile		
1025	1030	1035
Arg Asn Gly Ala Ser Phe Ser Tyr Cys Ser Lys Asp Arg		1040
1045	1050	

<210> 275
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 275
aattctccga acgtgtcacg ttt

<210> 276
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 276
aaggcacgaa gacgaacacu uctt

24

<210> 277
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 277
aactgaagac ctgaagacaa taa

23